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GenCore version 5.1.6

OM nucleic - nucleic search, using sw mode!

Run on: August 6, 2005, 22:19:03 ; Search time 3127.83 Seconds

(without alignments)
10826.558 Million cell updates/sec

Title: US-09-509-591-2

Perfect score: 5224 Sequence: 1 GAACTTAGAAATATGGG.....TTCCACCACTGCCATTACA 5224

Scoring table: IDENTITY NUC Gapop 10.0 , Gapext 1.0

Searched: 7297361 seqs, 3241162794 residues

Total number of hits satisfying chosen parameters:

14594722

Minimum DB seq length: 0

Maximum DB seq length: 20000000000

Post-processing: Minimum Match 0% Maximum Match 100%

Listing first 45 summaries

Database : Published Applications NA:*

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RESULT 1
US-09-509-883-2

Sequence 2, Application US/09898883
Patent No. US002016479A1

GENERAL INFORMATION:
APPLICANT: Little, Andrew
Lamparski, Henry
Schaur, Eric

TITLE OF INVENTION: ADENOVIRUS VECTORS SPECIFIC FOR CELLS
EXPRESSING APHA-PETROPROTEIN AND METHODS OF USE THEREOF
NUMBER OF SEQUENCES: 23
CORRESPONDENCE ADDRESS:
ADDRESSEE: MORRISON & FOERSTER
STREET: 755 PAGE MILL ROAD
CITY: PALO ALTO
STATE: CA
COUNTRY: USA
ZIP: 94304-1018

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
CURRENT APPLICATION DATA: #1.0, Version #1.30
APPLICATION NUMBER: US/09/898-883

FILING DATE: 02-Jul-2001

CLASSIFICATION: <Unknown>

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US/09/033,428

SUMMARIES

| Result No. | Score | Query Match | Length | DB ID | Description |
|------------|-------|-------------|--------|-------|------------------|
| 1 | 5224 | 100.0 | 5224 | 9 | US-09-898-883-2 |
| 2 | 5224 | 100.0 | 5224 | 10 | US-09-151-376-5 |
| 3 | 5224 | 100.0 | 5224 | 10 | US-09-151-376-45 |
| 4 | 5224 | 100.0 | 5224 | 10 | US-09-814-357-16 |
| 5 | 5224 | 100.0 | 5224 | 10 | US-09-814-351-16 |
| 6 | 5224 | 100.0 | 5224 | 15 | US-10-226-820-15 |
| 7 | 5224 | 100.0 | 5224 | 16 | US-10-139-089-5 |

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

| Result No. | Score | Query Match | Length | DB ID | Description |
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| 1 | 5224 | 100.0 | 5224 | 9 | Sequence 2, Appli |
| 2 | 5224 | 100.0 | 5224 | 10 | Sequence 5, Appli |
| 3 | 5224 | 100.0 | 5224 | 10 | Sequence 45, Appli |
| 4 | 5224 | 100.0 | 5224 | 10 | Sequence 16, Appli |
| 5 | 5224 | 100.0 | 5224 | 10 | Sequence 16, Appli |
| 6 | 5224 | 100.0 | 5224 | 15 | Sequence 15, Appli |
| 7 | 5224 | 100.0 | 5224 | 16 | Sequence 1, Appli |

FILING DATE: <Unknown>
ATTORNEY/AGENT INFORMATION:
NAME: POLIZZI, CATHERINE M.
REGISTRATION NUMBER: 40 130
REFERENCE/DOCKET NUMBER: 34802-30004 .00
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 813-5600
TELEFAX: (415) 494-0792
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 5244 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: Linear
SEQUENCE DESCRIPTION: SEQ ID NO: 2:
us-09-898-883-2

| Query Match | Score | DB | Length |
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| Best Local Similarity | 100.0% | 9 | 3224; |
| Matches 524: | Conservative | 0; | Pred. No. 0; |
| | Mismatches | 0; | Indels 0; |
| | Gaps | 0; | |
| Qy | 1 | GAATTCCTAGAAATATGGGGTAAATGCTGTTAACCCCATAGGT | 60 |
| Db | 1 | GAATTCCTAGAAATATGGGGTAAATGCTGTTAACCCCATAGGT | 60 |
| Qy | 61 | GAGATAGCATGGTTAACACATACATAGATAAAGG | 120 |
| Db | 61 | GAGATAGCATGGTTAACACATACATAGATAAAGG | 120 |
| Qy | 121 | AACAGACTATGGCTGGGACTTGAGATGCTGTCATAAACCTGGGTCTATCT | 180 |
| Db | 121 | AACAGACTATGGCTGGGACTTGAGATGCTGTCATAAACCTGGGTCTATCT | 180 |
| Qy | 181 | GTTCPATGGGGCTCTGGTTAACGCTTGCACAGGGTTCACTGACTCTTCTCCC | 240 |
| Db | 181 | GTTCPATGGGGCTCTGGTTAACGCTTGCACAGGGTTCACTGACTCTTCTCCC | 240 |
| Qy | 241 | AAGCCCAAGGACTCTGCTCTTCTATCTGGGGCTCTGGGTTGATACT | 300 |
| Db | 241 | AAGCCCAAGGACTCTGCTCTTCTATCTGGGGCTCTGGGTTGATACT | 300 |
| Qy | 301 | GAGAAATAAACATTCAATAATGTTCTGGGAGATGAGATGATGAGATGTCAT | 360 |
| Db | 301 | GAGAAATAAACATTCAATAATGTTCTGGGAGATGAGATGATGAGATGTCAT | 360 |
| Qy | 361 | TCAATTGTTCAATGTAATGAGATGAGATGAGATGAGATGAGATGAGATGTCAT | 420 |
| Db | 361 | TCAATTGTTCAATGTAATGAGATGAGATGAGATGAGATGAGATGAGATGTCAT | 420 |
| Qy | 421 | GAGGTGAGGGTGTACTATCATTATGAGATCTTATTTA | 480 |
| Db | 421 | GAGGTGAGGGTGTACTATCATTATGAGATCTTATTTA | 480 |
| Qy | 481 | TGCTTGTACAATGTTGTTGGGACACAGGATACAAAGATGTCCTGTAATT | 540 |
| Db | 481 | TGCTTGTACAATGTTGTTGGGACACAGGATACAAAGATGTCCTGTAATT | 540 |
| Qy | 541 | AAGAACTTAATGTTGAGGAAATTACATAGCTTCAAAATGACTGATAACCTAA | 600 |
| Db | 541 | AAGAACTTAATGTTGAGGAAATTACATAGCTTCAAAATGACTGATAACCTAA | 600 |
| Qy | 601 | ACAGGGTTCCATGAAATAATGTTAAAGGTTTAAAGGTGTCACAGGTGAGGCG | 660 |
| Db | 601 | ACAGGGTTCCATGAAATAATGTTAAAGGTTTAAAGGTGTCACAGGTGAGGCG | 660 |
| Qy | 661 | CTCTTCTCTGCTAGATGAGATGATGAACTTCAGGATAATTATGATAAGCT | 720 |
| Db | 661 | CTCTTCTCTGCTAGATGAGATGATGAACTTCAGGATAATTATGATAAGCT | 720 |
| Qy | 721 | TAAGGGTTGGSTACATTGTTCAACCATGTTGAGCTTCACTACATCAGGAAAC | 780 |
| Db | 721 | TAAGGGTTGGSTACATTGTTCAACCATGTTGAGCTTCACTACATCAGGAAAC | 780 |

Qy 781 ATTTCAACTAACCCACCACCCATTCCAAACAAAAATGAAAGAACTCAGAAACA 840
Db 781 ATTTCAACTAACCCACCACCCATTCCAAACAAAAATGAAAGAACTCAGAAACA 840
Qy 841 GTGAGATAAGGAGGAATTTCCTACAACTCCACAGTATAGCTAACCTGCTGAGAA 900
Db 841 GTGAGATAAGGAGGAATTTCCTACAAACCCACCGTATAGCTAACCTGCTGAGAA 900
Qy 901 GTATATATCTAAATTAACTACATGCPATAATGATAATAATTACTGTCATT 960
Db 901 GTATATATCTAAATTAACTACATGCPATAATGATAATAATTACTGTCATT 960
Qy 961 TTTATGTCATAAGCTACAGGGCATTTAGAAGATTTACCTTATATCAAATAA 1020
Db 961 TTAAATGTCATAACGGGATTAGAGATTAACTCCATTATATCAAATAA 1020
Qy 1021 ACTTGAGGGATAGTCATACACTAACTGTCATTTCAGTTGAACTGAGACT 1080
Db 1021 ACTTGAGGGATAGTCATACACTAACTGTCATTTCAGTTGAACTGAGACT 1080
Qy 1081 TTGCTGTCTACACTAACTGTCATTTCAGTTGAACTGCTGTTCTGTTGAGCT 1140
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Qy 1141 CTTCTTAATACCAAGGTCAGTTAACCTGTCATTTCAGTTGAACTGCTGTT 1200
Db 1141 CTTCTTAATACCAAGGTCAGTTAACCTGTCATTTCAGTTGAACTGCTGTT 1200
Qy 1201 TATGAAAGTAAACGGATTGAACTCATGTCATAAGCTAACAGCATGTCATGCTGTT 1260
Db 1201 TATGAAAGTAAACGGATTGAACTCATGTCATAAGCTAACAGCATGTCATGCTGTT 1260
Qy 1261 ACTTGGACTAACTTGAGCAATAACACATAGATACTGGATTGTTCTGTAGCAT 1320
Db 1261 ACTTGGACTAACTTGAGCAATAACACATAGATACTGGATTGTTCTGTAGCAT 1320
Qy 1321 ACAACACTCCTGTTCAAGCTCCTTAAAGCTCCTCTTAAAGCTGTCCTTCAT 1380
Db 1321 ACAACACTCCTGTTCAAGCTCCTCTTAAAGCTGTCCTCTTAAAGCTGTCCTTCAT 1380
Qy 1381 GGTTCTCTTCTGCTATGTTCTCACACTICAATGCTACATAACTGTC 1440
Db 1381 GGTTCTCTTCTGCTATGTTCTCACACTICAATGCTACATAACTGTC 1440
Qy 1441 TGCAAGCTTATGTTGATCTCTGCTAAATAATCTATCTGCTAACTGTC 1500
Db 1441 TGCAAGCTTATGTTGATCTCTGCTAAATAATCTATCTGCTAACTGTC 1500
Qy 1501 AGTAGTAACTCAAACTCACATCAACCTCCACTTCAACATA 1560
Db 1501 AGTAGTAACTCAAACTCACATCAACCTCCACTTCAACATA 1560
Qy 1561 CAAACGGGGAGTTGCTTCTAAATAATGTTCCCTGCTAGGACATAGGGTCC 1620
Db 1561 CAAACGGGGAGTTGCTTCTAAATAATGTTCCCTGCTAGGACATAGGGTCC 1620
Qy 1621 TCTTGTCCCTTAATACCTTAACTACTGCTAACTGCTAACTGCTAACTGCT 1740
Db 1621 TCTTGTCCCTTAATACCTTAACTACTGCTAACTGCTAACTGCTAACTGCT 1740
Qy 1741 GTTTGGTAAACTGCTCACTTTCACTTAACTAAACTAAACTGCTAACTGCT 1800
Db 1741 GTTTGGTAAACTGCTCACTTTCACTTAACTAAACTAAACTGCTAACTGCT 1800
Qy 1801 AGTACTAAGTCTTGTGACTTATCTCATCAGGAACTGCTTATCCAGGCCCTTAT 1860
Db 1801 AGTACTAAGTCTTGTGACTTATCTCATCAGGAACTGCTTATCCAGGCCCTTAT 1860

| | | | |
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| 4 | 4021 | GTCATGATCACAACTGAGGCACTCTCCATGTATTGTTCTGGCTGTTTCAAGTTA | 4 080 |
| Qy | 4 081 | ACAATATTATGAGCATATTCCATTTCAATAATTCTTCCAAAGTTATTTCAT | 4 140 |
| Db | 4 081 | ACAATATTATGAGCATATTCCATTTCAATAATTCTTCCAAAGTTATTTCAT | 4 140 |
| Qy | 4 141 | GGCTGATATACCCACTTTATGAAATGACCATTAATTATTCTGGCTGGGTTA | 4 200 |
| Db | 4 141 | GGCTGATATACCCACTTTATGAAATGACCATTAATTATTCTGGCTGGGTTA | 4 200 |
| Qy | 4 201 | TTTGATTTTATAATCTTACCTTTAGATAATGAAACCTGTGAAGCTTAAATACT | 4 260 |
| Db | 4 201 | TTTGATTTTATAATCTTACCTTTAGATAATGAAACCTGTGAAGCTTAAATACT | 4 260 |
| Qy | 4 261 | GGTGCCTGGTCTCAGATCTCCAGATTCCTGTTAATGTTCTGGTTACAGACTGCCA | 4 320 |
| Db | 4 261 | GGTGCCTGGTCTCAGATCTCCAGATTCCTGTTAATGTTCTGGTTACAGACTGCCA | 4 320 |
| Qy | 4 321 | TTGGGAAATTCAAAGCTCCCGACGTGATCTAATGTTAGCCAAATGGGAACCCCTG | 4 380 |
| Db | 4 321 | TTGGGAAATTCAAAGCTCCCGACGTGATCTAATGTTAGCCAAATGGGAACCCCTG | 4 380 |
| Qy | 4 381 | TAGACGGGATGATAGGGTAGCCACTCTAGCATTTAGTATTAACATCAT | 4 440 |
| Db | 4 381 | TAGACGGGATGATAGGGTAGCCACTCTAGCATTTAGTATTAACATCAT | 4 440 |
| Qy | 4 441 | CATCTTGAGTTGCTAATGTAATGCACTTATAAGAACATGTTG2AA | 4 500 |
| Db | 4 441 | CATCTTGAGTTGCTAATGTAATGCACTTATAAGAACATGTTG2AA | 4 500 |
| Qy | 4 501 | AAAAATTATTTAGGGCTTATTAGGGCTTATTAGGGCTTATTAGGGCTTATTAGG | 4 560 |
| Db | 4 501 | AAAAATTATTTAGGGCTTATTAGGGCTTATTAGGGCTTATTAGGGCTTATTAGG | 4 560 |
| Qy | 4 561 | ATACATGCATCAACTAAATCTTAATGTAACCTTTAGTGCATACATATTAGTGAAG | 4 620 |
| Db | 4 561 | ATACATGCATCAACTAAATCTTAATGTAACCTTTAGTGCATACATATTAGTGAAG | 4 620 |
| Qy | 4 621 | TGTTCTGAGCTAAAGCATGAGCTAAATTATCAACATGAAATTGAAATGTT | 4 680 |
| Db | 4 621 | TGTTCTGAGCTAAAGCATGAGCTAAATTATCAACATGAAATTGAAATGTT | 4 680 |
| Qy | 4 681 | ATTATTCTGCAACTTGGGACAGTCATCTCTGAAATTTTGTAATTGAGGATT | 4 740 |
| Db | 4 681 | ATTATTCTGCAACTTGGGACAGTCATCTCTGAAATTTTGTAATTGAGGATT | 4 740 |
| Qy | 4 741 | GTTATTTGCAAGAAGTGAAGGTCGAAATTGTCAGAACATTGCTGGCATA | 4 800 |
| Db | 4 741 | GTTATTTGCAAGAAGGTCGAAATTGTCAGAACATTGCTGGCATA | 4 800 |
| Qy | 4 801 | TGATAGGCATTAAATGTTAAAGGAAATTGTTAGTGAATTGCAACAACT | 4 860 |
| Db | 4 801 | TGATAGGCATTAAATGTTAAAGGAAATTGTTAGTGAATTGCAACAACT | 4 860 |
| Qy | 4 861 | GCTGCTTCTCTGGCTCATTAACCTTAACTGAAATTCTGAGGAAATTATTCTGCAACT | 4 920 |
| Db | 4 861 | GCTGCTTCTCTGGCTCATTAACCTTAACTGAAATTATTCTGCAACT | 4 920 |
| Qy | 4 921 | TAGGGCAACATGCTGCTGAAATTCTGAGGAAATTATTCTGAGGAAATTATTCTGCAACT | 4 980 |
| Db | 4 921 | TAGGGCAACATGCTGCTGAAATTCTGAGGAAATTATTCTGAGGAAATTATTCTGCAACT | 4 980 |
| Qy | 4 981 | AAAATAATAAGTTGCAAGTTTTCTGCTTTGAAATTCTGAGGAAATTATTCTGCAACT | 5 040 |
| Db | 4 981 | AAAATAATAAGTTGCAAGTTTTCTGCTTTGAAATTCTGAGGAAATTATTCTGCAACT | 5 040 |
| Qy | 5 041 | ATAAAATACAAATAACCGPATGCTGTTAAATTGAGGAAATTATTCTGCAACT | 5 100 |
| Db | 5 041 | ATAAAATACAAATAACCGPATGCTGTTAAATTGAGGAAATTATTCTGCAACT | 5 100 |
| Qy | 5 101 | GGAAATACCAATAAGTAACTGATATACCAACAAAGGTTACTAGTAAAGGCTTGCT | 5 160 |
| Db | 5 101 | GGAAATACCAATAAGTAACTGATATACCAACAAAGGTTACTAGTAAAGGCTTGCT | 5 160 |

| | | | | |
|----|------|---|--|--|
| Qy | 541 | AAGAAGTAAATGGTCACCGAAATTAACTAGCTTACAAATGACTATGATAACCCCAA 600 | | |
| Db | 541 | AAGAAGTAAATGGTCACCGAAATTAACTAGCTTACAAATGATAACCCCAA 600 | | |
| Qy | 601 | ACAGAGGTTCCATGAGAAATACTGAAAGGTTAAATAGTGTAAAGTGAGGG 660 | | |
| Db | 601 | ACAGAGGTTCCATGAGAAATACTGAAAGGTTAAATAGTGTAAAGTGAGGG 660 | | |
| Qy | 661 | CCTTCCTCTAGCTAGAGACTAACTCAGAAATACTTCAAGGATAATTGTAAGACT 720 | | |
| Db | 661 | CCTTCCTCTAGCTAGAGACTAACTCAGAAATACTTCAAGGATAATTGTAAGACT 720 | | |
| Qy | 721 | TAAGGGTTGGTACATTGTCAAGGATTGTGGAGAGGAGCTGAAATTGAAAAAC 780 | | |
| Db | 721 | TAAGGGTTGGTACATTGTCAAGGATTGTGGAGAGGAGCTGAAATTGAAAAAC 780 | | |
| Qy | 781 | ATTTCACTAACCAACCCAAATCCAACACAACCTATCTCAGAAA CA 840 | | |
| Db | 781 | ATTTCACTAACCAACCCAAATCCAACACAACCTATCTCAGAAA CA 840 | | |
| Qy | 841 | GTGAGATAAGAGGAAATTCTCACACCCCCACCTATACCTGCTGAGAA 900 | | |
| Db | 841 | GTGAGATAAGAGGAAATTCTCACACCCCCACCTATACCTGCTGAGAA 900 | | |
| Qy | 901 | GTTATATCTAAATTAACTAACATACTCATGCTTAATACTGCTATT 960 | | |
| Db | 901 | GTTATATCTAAATTAACTAACATACTCATGCTTAATACTGCTATT 960 | | |
| Qy | 961 | TTAATGCTTAAAGTACAGGATTATCCATTATATCAAAATA 1020 | | |
| Db | 961 | TTAATGCTTAAAGTACAGGATTATCCATTATATCAAAATA 1020 | | |
| Qy | 1021 | ACTTGAGGGTCTCATACAGCTTACATTGACCATAAGACATTGAAAT 1080 | | |
| Db | 1021 | ACTTGAGGGTCTCATACAGCTTACATTGACCATAAGACATTGAAAT 1080 | | |
| Qy | 1081 | TTGCCCTCATACAGCTTACATTGACCATAAGACATTGAAAT 1140 | | |
| Db | 1081 | TTGCCCTCATACAGCTTACATTGACCATAAGACATTGAAAT 1140 | | |
| Qy | 1141 | CTTCTTAATACCAAGTTCTGTTACTGTTCCATGTTGCTTCAGACT 1200 | | |
| Db | 1141 | CTTCTTAATACCAAGTTCTGTTACTGTTCCATGTTGCTTCAGACT 1200 | | |
| Qy | 1201 | TATGAAAAGTAAACGGATACTGAAATTCACTAAATGAAAGGCTATCTGT 1260 | | |
| Db | 1201 | TATGAAAAGTAAACGGATACTGAAATTCACTAAATGAAAGGCTATCTGT 1260 | | |
| Qy | 1261 | ACTTGGCTTAACACTTGAGGAAATAACACATAGTGGGATTGTTGTGTAGCAT 1320 | | |
| Db | 1261 | ACTTGGCTTAACACTTGAGGAAATAACACATAGTGGGATTGTTGTGTAGCAT 1320 | | |
| Qy | 1321 | ACAAACTCTGGTTCAAGCTCTTCTTCTGTTGAAATTGGCTTCAT 1380 | | |
| Db | 1321 | ACAAACTCTGGTTCAAGCTCTTCTTCTGTTGAAATTGGCTTCAT 1380 | | |
| Qy | 1381 | GTTTCTCTTCTACGTCTTCAACCTACATGCTCAATACTGTC 1440 | | |
| Db | 1381 | GTTTCTCTTCTACGTCTTCAACCTACATGCTCAATACTGTC 1440 | | |
| Qy | 1441 | TGCAAGGTATGATTAATCTGCTTAACCTGCTTCAAGGATAAAA 1500 | | |
| Db | 1441 | TGCAAGGTATGATTAATCTGCTTAACCTGCTTCAAGGATAAAA 1500 | | |
| Qy | 1501 | AGTAGTATCAATGCACTAACGCTTCCACTTGTGTTAAAGCTTCAACATA 1560 | | |
| Db | 1501 | AGTAGTATCAATGCACTAACGCTTCCACTTGTGTTAAAGCTTCAACATA 1560 | | |
| Qy | 1561 | CAAACCGGGAGTTGCTGAAATTGTGTCCTGAGACATAGGTCC 1620 | | |
| Db | 1561 | CAAACCGGGAGTTGCTGAAATTGTGTCCTGAGACATAGGTCC 1620 | | |
| Qy | 1621 | TCTTGTCTCTTAAATCTAACTTGAAGCTGTCTCATCCACCTATGGGAGATG 1680 | | |
| Db | 1621 | TCTTGTCTCTTAAATCTAACTTGAAGCTGTCTCATCCACCTATGGGAGATG 1680 | | |
| Qy | 1681 | AGAGTGAAGGGAGCTGTTAACTGATACTGCTTAACTAACTGAACTGAGACT 1740 | | |
| Db | 1681 | AGAGTGAAGGGAGCTGTTAACTGATACTGCTTAACTAACTGAACTGAGACT 1740 | | |
| Qy | 1741 | GTTCGGTTAACTGTCACTTATCTTAACTAACTGAACTGACTT 1800 | | |
| Db | 1741 | GTTCGGTTAACTGTCACTTATCTTAACTAACTGAACTGACTT 1800 | | |
| Qy | 1801 | AGTTACTAAGCTCTGACTTTGACTTCTCATACACTCATGCTTATCCAGGCCACTT 1860 | | |
| Db | 1801 | AGTTACTAAGCTCTGACTTTGACTTCTCATACACTCATGCTTATCCAGGCCACTT 1860 | | |
| Qy | 1861 | TTCAGTGTATTATTGGAAACCTCTTCAACTGTTCTCTTCTATCATGCTTATCCCTT 1920 | | |
| Db | 1861 | TTCAGTGTATTATTGGAAACCTCTTCAACTGTTCTCTTCTATCATGCTTATCCCTT 1920 | | |
| Qy | 1921 | TGAAACRAAAAGGAGAGCTTCAAAATACTGTTTATGTTTATAGTCCCTTGTGT 1980 | | |
| Db | 1921 | TGAAACRAAAAGGAGAGCTTCAAAATACTGTTTATGTTTATAGTCCCTTGTGT 1980 | | |
| Qy | 1981 | CTATATACTGCCCAGAAGGGATTATAACTCCATTAAAGCTTCAACTACATATA 2040 | | |
| Db | 1981 | CTATATACTGCCCAGAAGGGATTATAACTCCATTAAAGCTTCAACTACATATA 2040 | | |
| Qy | 2041 | TGCCAATCTTGCAGAAATTCCAAATATCPAGTATTTCTTAACTTAACTTAACTTGTGCTC 2100 | | |
| Db | 2041 | TGCCAATCTTGCAGAAATTCCAAATATCPAGTATTTCTTAACTTAACTTAACTTGTGCTC 2100 | | |
| Qy | 2101 | TTCAAAACTGCTATTCTCTCATCCCTTAAGTGTCACTCCCTAACCTGGTTGTT 2160 | | |
| Db | 2101 | TTCAAAACTGCTATTCTCTCATCCCTTAAGTGTCACTCCCTAACCTGGTTGTT 2160 | | |
| Qy | 2161 | TTCCACCACTTACATTCTGATTCATTCTCTCATCCCTAACCTGGTTGTT 2220 | | |
| Db | 2161 | TTCCACCACTTACATTCTGATTCATTCTCTCATCCCTAACCTGGTTGTT 2220 | | |
| Qy | 2221 | TCTAATTCTCTCATGCTCCATGAAAGTTTACCTTCTCTTACCTGGTTGCTC 2280 | | |
| Db | 2221 | TCTAATTCTCTCATGCTCCATGAAAGTTTACCTTCTCTTACCTGGTTGCTC 2280 | | |
| Qy | 2281 | CCTCCAAAGATGTCTGAGTTCTCTTCACTTCTTCACTTCACTACACCA 2340 | | |
| Db | 2281 | CCTCCAAAGATGTCTGAGTTCTCTTCACTTCTTCACTTCACTACACCA 2340 | | |
| Qy | 2341 | TGTTGTGATTACTGATACTATTGTTGTTCTGATTAGCTAACAGAGC 2400 | | |
| Db | 2341 | TGTTGTGATTACTGATACTATTGTTGTTCTGATTAGCTAACAGAGC 2400 | | |
| Qy | 2401 | TACATGTGCTCTCTTGTGCTTAACTGAAATTCTCATCCATC 2460 | | |
| Db | 2401 | TACATGTGCTCTCTTGTGCTTAACTGAAATTCTCATCCATC 2460 | | |
| Qy | 2461 | GACTTAACTTTTATGAAATTAAACCCCATCTPATGAGTGTGCTACTTGTGCA 2520 | | |
| Db | 2461 | GACTTAACTTTTATGAAATTAAACCCCATCTPATGAGTGTGCTACTTGTGCA 2520 | | |
| Qy | 2521 | AGACCCGGTTCTGAGSCATTATATTGTTTATCTCATTAACCATGAAAG 2580 | | |
| Db | 2521 | AGACCCGGTTCTGAGSCATTATATTGTTTATCTCATTAACCATGAAAG 2580 | | |
| Qy | 2581 | GAGGACTACTACATCTCCATTCTGTTAACTGAAATTCAACCACTTCCCAACT 2640 | | |
| Db | 2581 | GAGGACTACTACATCTCCATTCTGTTAACTGAAATTCAACCACTTCCCAACT 2640 | | |
| Qy | 2641 | AACCTACCCAAAGTGTGACTCTGAGCTTCCCAACT 2700 | | |
| Db | 2641 | AACCTACCCAAAGTGTGACTCTGAGCTTCCCAACT 2700 | | |
| Qy | 2701 | TTACCTGATTAACTCTGCTTACTGCTCATGGCATGGAGAT 2760 | | |

4 921 TAGGACAAGTCATCTTGAAATTCTTGATTTCTGAGGTATTGGTATATTTGC 4 980
 Qy ;
 Db 4 921 TAGGACAAGTCATCTTGAAATTCTTGATTTCTGAGGTATTGGTATATTTGC 4 980
 Qy ;
 Db 4 981 AAAATAAAATAAAGTTGCAAGTTTCTGCCCAAAGACTCTGCTCTGAC 5 040
 Qy ;
 Db 4 981 AAAATAAAATAAAGTTGCAAGTTTCTGCCCAAAGACTCTGCTCTGAC 5 040
 Qy ;
 Db 5041 ATAAATAACAATAACCGCTATGCCTTAATTGGCAAATGTCACCTAA 5 100
 Qy ;
 Db 5041 ATAAATAACAATAACCGCTATGCCTTAATTGGCAAATGTCACCTAA 5 100
 Qy ;
 Db 5101 GGAATTACCATTAAGTAAAGATATAACCAAAAGGTACTACTAACGGATATGGCT 5 160
 Qy ;
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 Qy ;
 Db 5161 GAAAGAGTATAAAAAGAATTTCAGCATGATTTCACATTTGCTTCCACCACTGCCATT 5 220
 Qy ;
 Db 5161 GAAAGAGTATAAAAAGAATTTCAGCATGATTTCACATTTGCTTCCACCACTGCCATT 5 220
 Qy 5221 AACAA 5224
 Qy ;
 Db 5221 AACAA 5224
 Qy ;
 Db 5221 AACAA 5224

RESULT 3

US-09-151-376-45
 ; Sequence 45, Application US/09151376
 ; Publication No. US20030044383A1
 ; GENERAL INFORMATION:
 ; APPLICANT: Henderson, D.R.
 ; ATTORNEY: Schaur, E.R.
 ; TITLE OF INVENTION: TISSUE SPECIFIC VIRAL VECTORS
 ; FILE REFERENCE: 34802000021
 ; CURRENT APPLICATION NUMBER: US/09/151,376
 ; CURRENT FILING DATE: 1998-09-10
 ; EARLIER APPLICATION NUMBER: 08/669,753
 ; EARLIER FILING DATE: 1996-06-26
 ; EARLIER APPLICATION NUMBER: 08/495,034
 ; EARLIER FILING DATE: 1995-06-27
 ; NUMBER OF SEQ ID NOS: 71
 ; SOFTWARE: PatentIn Ver. 2.0
 ; SEQ ID NO: 45
 ; LENGTH: 5224
 ; TYPE: DNA
 ; ORGANISM: Unknown
 ; FEATURE:
 ; OTHER INFORMATION: Description of Unknown Organism: unknown
 US-09-151-376-45

Query Match Score 5224; DB 10; Length 5224;
 Best Local Similarity 100.0%; Pred. No. 0;
 Matches 5224; Mi matches 0; Indels 0; Gaps 0;

| | |
|--------|---|
| Qy 1 | GAATCTTGAATAATGGGGTAAATGTSCTTCAACACATCATTCAAGATTAAGG 120 |
| Db 1 | GAATCTTGAATAATGGGGTAAATGTSCTTCAACACATCATTCAAGATTAAGG 120 |
| Qy 61 | GAGATAAGCATTGGTTAAATGTSCTTCAACACATCATTCAAGATTAAGG 120 |
| Db 61 | GAGATAAGCATTGGGGTAAATGTSCTTCAACACATCATTCAAGATTAAGG 120 |
| Qy 121 | AACAGACATATGGCTGGAGGACCTTGAGATGCTGCTCATACACTGGGTGTATCT 180 |
| Db 121 | AACAGACATATGGGGTGGAGGACCTTGAGATGCTGCTCATACACTGGGTGTATCT 180 |
| Qy 181 | GTCTATGGGCTGTGTTAAGCTGGAAACTTGCAACGGGTCACTGACTTTCTCCC 240 |
| Db 181 | GTCTATGGGCTGTGTTAAGCTGGAAACTTGCAACGGGTCACTGACTTTCTCCC 240 |
| Qy 241 | AAGCCCAAGCTGCTACTGCTCTTCAATCTGTTGGGCTCTGGGCTGAAATATCT 300 |
| Db 241 | AAGCCCAAGCTGCTACTGCTCTTCAATCTGTTGGGCTCTGGGCTGAAATATCT 300 |

| | | |
|------|--|-------|
| 3541 | TGGAATTCAAAGGGTAGACAGGTTGGTGTACTCTTCTGGAGAGTACGAGAACGAGAAG | 3 600 |
| 3601 | GCCCTTTGTTAACHTGTCTTAGGAGCAAACCTTCTGGAGAACGAGAACGCTAGCGAA | 3 660 |
| 3601 | GCCCTTTGTTAACHTGTCTTAGGAGCAAACCTTCTGGAGAACGAGAACGCTAGCGAA | 3 660 |
| 3661 | AATGGCAAAACTACAATGAATGGAATTGTCTGATTAGCTTGAGACCTGTGTT | 3 720 |
| 3661 | AATGGCAAAACTACAATGAATGGAATTGTCTGATTAGCTTGAGACCTGTGTT | 3 720 |
| 3721 | TATACTATGATAAACTTTGPAATTCTGGAAAGTGTACTGACGGTAACCCCTTTGTGTT | 3 780 |
| 3721 | TATACTATGATAAACTTTGPAATTCTGGAAAGTGTACTGACGGTAACCCCTTTGTGTT | 3 780 |
| 3781 | TTAATGTTGTCGCCCTAGTAGCTTGCAAGTGTCTGATCTTAAAGTACTGTACTTGTGTT | 3 840 |
| 3781 | TTAATGTTGTCGCCCTAGTAGCTTGCAAGTGTCTGATCTTAAAGTACTGTACTTGTGTT | 3 840 |
| 3841 | TTTAAAATTTATGTTAAATGGATAGTGTCTCTTCACTGAAAGAAGTTTGAAAGAG | 3 900 |
| 3841 | TTTAAAATTTATGTTAAATGGATAGTGTCTCTTCACTGAAAGAAGTTTGAAAGAG | 3 900 |
| 3901 | AGATAGAAATTCAACTTCACTTCAACCATTAGAGAAACCCAATGTTAAACACTTGTGTT | 3 960 |
| 3901 | AGATAGAAATTCAACTTCACTTCAACCATTAGAGAAACCCAATGTTAAACACTTGTGTT | 3 960 |
| 3961 | TCCATTATTCTGCTTTATTCAACATTTTT2AGGGTGGGGAAATACAGGGAG | 4 020 |
| 3961 | TCCATTATTCTGCTTTATTCAACATTTTT2AGGGTGGGGAAATACAGGGAG | 4 020 |
| 4021 | GTACAAATGATAACAATGAGGAAACTCTCCATGATTGTTGCTGTGTTAGCTTA | 4 080 |
| 4021 | GTACAAATGATAACAATGAGGAACTCTCCATGATTGTTGCTGTGTTAGCTTA | 4 080 |
| 4081 | ACATAATTATGACCATATTCCATTTCATTAATAATTCTCACAAGTTATTGTAT | 4 140 |
| 4081 | ACATAATTATGACCATATTCCATTTCATTAATAATTCTCACAAGTTATTGTAT | 4 140 |
| 4141 | GGCTGTATATCACCCCTACTTTATGAAATGACCATATAATTCTCTGTTGCGTTA | 4 200 |
| 4141 | GGCTGTATATCACCCCTACTTTATGAAATGACCATATAATTCTCTGTTGCGTTA | 4 200 |
| 4201 | TTTGATTTATAATCTTACCTTATGAAATGAAACCTGTGAACTTTGAAATACT | 4 260 |
| 4201 | TTTGATTTATAATCTTACCTTATGAAATGAAACCTGTGAACTTTGAAATACT | 4 260 |
| 4261 | GGTCCCTGGCTCACTCCACAGATTCGTTGACCTTGCTGCTGTTACGACTGGCA | 4 320 |
| 4261 | GGTCCCTGGCTCACTCCACAGATTCGTTGACCTTGCTGCTGTTACGACTGGCA | 4 320 |
| 4321 | TTGGAAATTCAAAGTCCCCAGTGTGATCTTAATGTTAGCCAAAGATCGGGAAACCTTG | 4 380 |
| 4321 | TTGGAAATTCAAAGTCCCCAGTGTGATCTTAATGTTAGCCAAAGATCGGGAAACCTTG | 4 380 |
| 4381 | TAGACGGGATGATGAGGTGAGCCACCTTAGCATCCATTAATGTTAACATCAT | 4 440 |
| 4381 | TAGACGGGATGATGAGGTGAGCCACCTTAGCATCCATTAATGTTAACATCAT | 4 440 |
| 4441 | CATCTTGACTTGTAAAGTGTGATGACCTGACCCACCTTATAAGACACATGTCAAA | 4 500 |
| 4441 | CATCTTGACTTGTAAAGTGTGATGACCTGACCCACCTTATAAGACACATGTCAAA | 4 500 |
| 4501 | TAAAATTATGAGCTTGTGTTTATAGGGCTCTAAGTTCTAGCTTGCTGTTCTAGTGTAGCC | 4 560 |
| 4501 | TAAAATTATGAGCTTGTGTTTATAGGGCTCTAGCTGCTGTTCTAGTGTAGCC | 4 560 |
| 4561 | ATACATCGATCAACTAAATCTTAAAGTACCTTGTACATATAAGCTAATGAAAG | 4 620 |
| 4561 | ATACATCGATCAACTAAATCTTAAAGTACCTTGTACATATAAGCTAATGAAAG | 4 620 |
| 4621 | TGTTCTGAGCTAAACAACTGATGTTGACAGCTTAACTGATGTTGAAATGAAATT | 4 680 |
| 4621 | TGTTCTGAGCTAAACAACTGATGTTGACAGCTTAACTGATGTTGAAATGAAATT | 4 680 |

RESULT 4

US-09-814-357-16
Sequence 16, Application US/09814357
Publication No. US20030068307A1

GENERAL INFORMATION:

APPLICANT: Yu, De-Chao
APPLICANT: Chen, Yu
APPLICANT: Henderson, Daniel R.

TITLE OF INVENTION: METHODS OF TREATMENT WITH COMBINATION CHEMOTHERAPY AND FILE REFERENCE: 349022001600

CURRENT APPLICATION NUMBER: US/09/8114357
CURRENT FILING DATE: 2001-10-15
PRIOR APPLICATION NUMBER: 60/192, 015
PRIOR FILING DATE: 2000-03-24

NUMBER OF SEQ ID NOS: 35
SOFTWARE: FastSEQ for Windows Version SEQ ID NO 16
LENGTH: 5224

TYPE: DNA
ORGANISM: Artificial sequence
FEATURE:
; OTHER INFORMATION: AlphaFP-TRE

US-09-814-357-16

Query Match 100 %
Best Local Similarity 100 %
Matches 5224; Conservative 0;
MFE 1

1 GAAATTCTAGAAATGGGGTAGG

| | | | |
|----|------|--|------|
| Qy | 2221 | TCTAATTTCAGTCCTCATGAGATGTACTTCCAGGAAGCCTTATCTGACC | 2280 |
| Dy | 2221 | TCTAATTTCAGTCCTCATGAGATGTACTTCCAGGAAGCCTTATCTGACC | 2280 |
| Db | 2281 | CCTCCAAGATGTCATAGTCCTCTCTTCACTTCACTAACAGCTTACACCA | 2340 |
| Dy | 2281 | CCTCCAAGATGTCATAGTCCTCTCTTCACTTCACTAACAGCTTACACCA | 2340 |
| Db | 2341 | TGGTGATTAATCTGATCATCTATGTCGTTCTGTTAGGGAGTAGCTCACAGGC | 2400 |
| Dy | 2341 | TGGTGATTAATCTGATCATCTATGTCGTTCTGTTAGGGAGTAGCTCACAGGC | 2400 |
| Db | 2341 | TGGTGATTAATCTGATCATCTATGTCGTTCTGTTAGGGAGTAGCTCACAGGC | 2400 |
| Dy | 2401 | TACATGGTCTGTCGTTCTGTTAGGGAGTAGCTCACAGGC | 2460 |
| Db | 2401 | TACATGGTCTGTCGTTCTGTTAGGGAGTAGCTCACAGGC | 2460 |
| Dy | 2461 | GACTTACATTTTATGAAATAAACCCATCATATCGAGTGCACCTTGCGA | 2520 |
| Db | 2461 | GACTTACATTTTATGAAATAAACCCATCATATCGAGTGCACCTTGCGA | 2520 |
| Dy | 2521 | AGACCCGGTCTGGGGATTTATTTTCACTTACCATGAAAG | 2580 |
| Db | 2521 | AGACCCGGTCTGGGGATTTATTTTCACTTACCATGAAAG | 2580 |
| Dy | 2581 | GAGGTACTATCACTATCCTTATTTTATGTTATTGATTAAGGAAATGAAATT | 2640 |
| Db | 2581 | GAGGTACTATCACTATCCTTATTTTATGTTATTGATTAAGGAAATGAAATT | 2640 |
| Dy | 2641 | AACCTACCCAAAGTCATGAACTTAAACCAATTAAGGAAATGAAATT | 2700 |
| Db | 2641 | AACCTACCCAAAGTCATGAACTTAAACCAATTAAGGAAATGAAATT | 2700 |
| Dy | 2701 | TTACGTTGATTAATCTGTCATCTGCTCTGATGAACTATGCAAT | 2760 |
| Db | 2701 | TTACGTTGATTAATCTGTCATCTGCTCTGATGAACTATGCAAT | 2760 |
| Dy | 2761 | CTGCTCGTAAGGCAAGTATGCAAGCAGGATGACACAAAACCGATTAAT | 2820 |
| Db | 2761 | CTGCTCGTAAGGCAAGTATGCAAGCAGGATGACACAAAACCGATTAAT | 2820 |
| Dy | 2821 | CAGAGGAAATTCACAGGAACTTCAACCGTAACTTCACTGAGTGTAGTC | 2880 |
| Db | 2821 | CAGAGGAAATTCACAGGAACTTCAACCGTAACTTCACTGAGTGTAGTC | 2880 |
| Dy | 2881 | GCATGCGACATCTTGATGAACTGGGGTGGATGACACAAAACCGATATA | 2940 |
| Db | 2881 | GCATGCGACATCTTGATGAACTGGGGTGGATGACACAAAACCGATATA | 2940 |
| Dy | 2941 | TTATTTGTTGTTGTTGTTGTTGTTGTTGTTGTTGTTGTTGTTGTTGTTG | 3000 |
| Db | 2941 | TTATTTGTTGTTGTTGTTGTTGTTGTTGTTGTTGTTGTTGTTGTTGTTG | 3000 |
| Dy | 3001 | TTACTAAGGCTTACCATTTGGCGATTTTACATTGTCCTCTATCTTGTG | 3060 |
| Db | 3001 | TTACTAAGGCTTACCATTTGGCGATTTTACATTGTCCTCTATCTTGTG | 3060 |
| Dy | 3061 | AGATGATGCAATTGGGATTAATCTCATCTATCATCTATCATCTATCATC | 3120 |
| Db | 3061 | AGATGATGCAATTGGGATTAATCTCATCTATCATCTATCATCTATCATC | 3120 |
| Dy | 3121 | TATATCTATCATCTATCATCTATCATCTATCATCTATCATCTATCATC | 3180 |
| Db | 3121 | TATATCTATCATCTATCATCTATCATCTATCATCTATCATCTATCATC | 3180 |
| Dy | 3181 | AATAATGTTCTCTTAAAGTTAAACCCATGTTGAGCTTAACTTCACTTCACT | 3240 |
| Db | 3181 | AATAATGTTCTCTTAAAGTTAAACCCATGTTGAGCTTAACTTCACTTCACT | 3240 |
| Dy | 3241 | GTAAACCCCTTCTCTCATATGTTGAAATGTTGAACTTCACTTCACT | 3300 |
| Db | 3241 | GTAAACCCCTTCTCTCATATGTTGAAATGTTGAACTTCACTTCACT | 3300 |

PRIOR APPLICATION NUMBER: 60/192,156
 PRIORITY FILING DATE: 2000-03-24
 NUMBER OF SEQ ID NOS: 35
 SOFTWARE: FastSEQ for Windows Version 4.0
 SEQ ID NO 16
 LENGTH: 5224
 TYPE: DNA
 ORGANISM: Artificial Sequence
 FEATURE:
 OTHER INFORMATION: AlphaSP-TRE
 US-09-814-351-16

| Query Match | Score 5224; | DB 10; | Length 5224; |
|-----------------------|---|---------------|--------------|
| Best Local Similarity | 100.0%; | Pred. No. 0; | |
| Matches 5224; | Conservative 0; | Mismatches 0; | Gaps 0 |
| Qy | 1 GAATCTTACAAATTGGGGTAGGGGCTGGGTAAATTCTGTTTCACCCCATAGGT 60 | | |
| Db | 1 GAATCTTACAAATTGGGGTAGGGGCTGGGTAAATTCTGTTTCACCCCATAGGT 60 | | |
| Qy | 61 GAGATAAGCATGGCTGGTTAATGCGCTTCACACACATCACATTTCATAGAACATTAGG 120 | | |
| Db | 61 GAGATAAGCATGGCTGGTTAATGCGCTTCACACACATCACATTTCATAGAACATTAGG 120 | | |
| Qy | 121 AACAGACTATGGCTGGGACTCTGGGATGTCGTTCTCATACACTTGEGGTTSTATCT 180 | | |
| Db | 121 AACAGACTATGGCTGGGACTCTGGGATGTCGTTCTCATACACTTGEGGTTSTATCT 180 | | |
| Qy | 181 GTTCTATGGGCCTGTTTAAGCTTGGCACAATTGGAAACGGTTACTGACTTTCCCC 240 | | |
| Db | 181 GTTCTATGGGCCTGTTTAAGCTTGGCACAATTGGAAACGGTTACTGACTTTCCCC 240 | | |
| Qy | 241 AAGCCAACTGACTCTCCCTCTGGCCCTCGGGCTGTGTTGAACTATCT 300 | | |
| Db | 241 AAGCCAACTGACTCTCCCTCTGGCCCTCGGGCTGTGTTGAACTATCT 300 | | |
| Qy | 301 GAGAAATAAAACTTCAATAATGTTCTGCTGGAGATGAGTATGAGATGTTCTGCAT 360 | | |
| Db | 301 GAGAAATAAAACTTCAATAATGTTCTGCTGGAGATGAGTATGAGATGTTCTGCAT 360 | | |
| Qy | 361 TCATTGTATCAAATGAAATTAGCTTGTAACTTCAAACTATCT 420 | | |
| Db | 361 TCATTGTATCAAATGAAATTAGCTTGTAACTTCAAACTATCT 420 | | |
| Qy | 421 GAGGTAGGGCTGGACTATTCAATTCTTCAATTAAAGATACTTCTTATTATTA 480 | | |
| Db | 421 GAGGTAGGGCTGGACTATTCAATTCTTCAATTAAAGATACTTCTTATTATTA 480 | | |
| Qy | 481 TGCCTGTGACAAATTGGGCTGGGACCAAGGAAATCACAAGGTGAGCTTGTGATT 540 | | |
| Db | 481 TGCCTGTGACAAATTGGGCTGGGACCAAGGAAATCACAAGGTGAGCTTGTGATT 540 | | |
| Qy | 541 AAGAGTTATGGTCAGGAATTATCATAGCTTACAATTGACTATGATAACATCAA 600 | | |
| Db | 541 AAGAGTTATGGTCAGGAATTATCATAGCTTACAATTGACTATGATAACATCAA 600 | | |
| Qy | 601 ACAAGAGTTCAATTGTCAGGAAATTAATCTGAAAGGTTAAATAGTGTCAAGGTGGAGGG 660 | | |
| Db | 601 ACAAGAGTTCAATTGTCAGGAAATTAATCTGAAAGGTTAAATAGTGTCAAGGTGGAGGG 660 | | |
| Qy | 661 CTCCTCTCTAGCTAGAGACTAAATCGAAATACATTGAGTATACTGAAATGACCT 720 | | |
| Db | 661 CTCCTCTCTAGCTAGAGACTAAATCGAAATACATTGAGTATACTGAAATGACCT 720 | | |
| Qy | 721 TAAGGTTGGGTACCTTGTCAAGCATGATGGAGAAGGAGCTGAAATTGAAAC 780 | | |
| Db | 721 TAAGGTTGGGTACCTTGTCAAGCATGATGGAGAAGGAGCTGAAATTGAAAC 780 | | |
| Qy | 781 ATTTCACAACTAACCAACCCAACTTCAGCTTGTGTTCTCAGAAACA 840 | | |
| Db | 781 ATTTCACAACTAACCAACCCAACTTCAGCTTGTGTTCTCAGAAACA 840 | | |
| Qy | 841 GTGAGGATAAGAGAGGAATTTCCTACACCCACGTATAGCTAACCTGCTCTAGAAAGAA 900 | | |
| Db | 841 GTGAGGATAAGAGAGGAATTTCCTACACCCACGTATAGCTAACCTGCTCTAGAAAGAA 900 | | |

RESULT 5
 US-09-814-351-16
 Sequence 16, Application US/0914351
 Title of Invention: CELL-SPECIFIC ADENOVIRUS VECTORS
 Comprising an Internal Ribosome Entry Site
 General Information:
 Current Filing Date: 2000-03-21
 Current Application Number: US/09/814,351
 Applicant: Yu, De-Chao
 Applicant: Li, Yuanhao
 Applicant: Henderson, Daniel R.
 Title of Invention: CELL-SPECIFIC ADENOVIRUS VECTORS
 Comprising an Internal Ribosome Entry Site
 File Reference: 3480220001700

| | | | | |
|----|------|---|------|--|
| Qy | 3061 | AGATGATCAGATTGGATTACTTGGCTTGAAGATGATAATCTACATCPATATC 3120 | 4141 | GGCTATATCACCCACTTTATGACCATTAATTCTGTGGGGTA 4200 |
| Db | 3061 | AGATGATCAGATTGGATTACTTGGCTTGAAGATGATAATCTACATCPATATC 3120 | 4141 | GGCTATATCACCCACTTTATGACCATTAATTCTGTGGGGTA 4200 |
| Qy | 3121 | TATATCTATCTATCTATCTATCTATCTATCTATCTATCTATCTATCTATCT 3180 | 4201 | TTCGTTTAAATACTTACCTTACGAAATATGAAACACCTGTGAAGCTT 4260 |
| Db | 3121 | TATATCTATCTATCTATCTATCTATCTATCTATCTATCTATCTATCTATCT 3180 | 4201 | TTCGTTTAAATACTTACCTTACGAAATATGAAACACCTGTGAAGCTT 4260 |
| Qy | 3181 | AAATATGTTTGTAAAGTTAAAGATTCAAGCTTTAGAAATCTGGATTGCCAAT 3240 | 4261 | GGTGTGGGTCTAACCTCCACATTCTGATTAACCTTGTTACAGACTGGCA 4320 |
| Db | 3181 | AAATATGTTTGTAAAGTTAAAGATTCAAGCTTTAGAAATCTGGATTGCCAAT 3240 | 4261 | GGTGTGGGTCTAACCTCCACATTCTGATTAACCTTGTTACAGACTGGCA 4320 |
| Qy | 3241 | GTAACCCTTCTCPACATTAACCCATGTTGGAAACAATACATTATTCTTCATC 3300 | 4321 | TTGGCAATTCAAAGTTCCCCTGGATCTTCAAGTGTAGCCAAAGTCCTTG 4380 |
| Db | 3241 | GTAACCCTTCTCPACATTAACCCATGTTGGAAACAATACATTATTCTTCATC 3300 | 4321 | TTGGCAATTCAAAGTTCCCCTGGATCTTCAAGTGTAGCCAAAGTCCTTG 4380 |
| Qy | 3301 | AAATGTTGCTCAAGTCTGGCTATGAACTGAACTGTTGGATATTTCGCC 3360 | 4381 | TAGACAGGGATGATGATGGCTGACCTCTGATCATTCTGATTAACATCAT 4440 |
| Db | 3301 | AAATGTTGCTCAAGTCTGGCTATGAACTGAACTGTTGGATATTTCGCC 3360 | 4381 | TAGACAGGGATGATGATGGCTGACCTCTGATCATTCTGATTAACATCAT 4440 |
| Qy | 3361 | CATGCTGGGAACTTGGCTTCAAAACTCTATTTCAGTCTTCATAACT 3420 | 4441 | CATCTGAGTTGCTAAGTGTGAACTGACCTTAACTTAAAGACATGTGCCAA 4500 |
| Db | 3361 | CATGCTGGGAACTTGGCTTCAAAACTCTATTTCAGTCTTCATAACT 3420 | 4441 | CATCTGAGTTGCTAAGTGTGAACTGACCTTAACTTAAAGACATGTGCCAA 4500 |
| Qy | 3421 | AATACTCTCACTATTGCTTCACTTCTTCATCAAAACTTGGCTTCATA 3480 | 4501 | TAAAATTATTATGAACTTGGTTTATTAGGCTTGTGCTTAACTTCTATGTAGCC 4560 |
| Db | 3421 | AATACTCTCACTATTGCTTCACTTCTTCATCAAAACTTGGCTTCATA 3480 | 4501 | TAAAATTATTATGAACTTCTGATTAACCTTAAAGTGTGAACTTAAAGAAAG 4620 |
| Qy | 3481 | TATTGAGTAAGTCCCTCTGAGAAAGTAGAGTAGAGAAACTCTGCACTTGTAAATCTACCC 3540 | 4561 | ATACATTCGCTAACTTGGCAAACTGATCTCTGTTGACTTTGTTGCTTAAG 4620 |
| Db | 3481 | TATTGAGTAAGTCCCTCTGAGAAAGTAGAGTAGAGAAACTCTGCACTTGTAAATCTACCC 3540 | 4561 | ATACATTCGCTAACTTGGCAAACTGATCTCTGTTGACTTTGTTGCTTAAG 4620 |
| Qy | 3541 | TGGAATCAACGGATAGCAAGGGATTGGCTTCTGAGAGTAGTGTGAG 3600 | 4621 | TGTTCTGAGCTAACANTGACCATTAATTCTGAGCTTAACTTGTGAAATT 4680 |
| Db | 3541 | TGGAATCAACGGATAGCAAGGGATTGGCTTCTGAGAGTAGTGTGAG 3600 | 4621 | TGTTCTGAGCTAACANTGACCATTAATTCTGAGCTTAACTTGTGAAATT 4680 |
| Qy | 3601 | GCCTGTTTGTTAACATGTTCTTACGGAAACCTTACGGAGACACGGCATAGCGAA 3660 | 4681 | ATTATTCGCAACTTGGCAAACTGATCTCTGTTGATTTCTGATGTTT 4740 |
| Db | 3601 | GCCTGTTTGTTAACATGTTCTTACGGAAACCTTACGGAGACACGGCATAGCGAA 3660 | 4681 | ATTATTCGCAACTTGGCAAACTGATCTCTGTTGATTTCTGATGTTT 4740 |
| Qy | 3661 | AATGGACAAAAACTAACAAATGAACTGGAAATTGACTGATTAGCATGAA 3720 | 4741 | GTATATTGCAAGCATGAGAGTCGAACATGCTGTGCTGGCTGGCATA 4800 |
| Db | 3661 | AATGGACAAAAACTAACAAATGAACTGGAAATTGACTGATTAGCATGAA 3720 | 4741 | GTATATTGCAAGCATGAGAGTCGAACATGCTGTGCTGGCTGGCATA 4800 |
| Qy | 3721 | TATACTATGATAAAGTTGTTATGCTGGAAAGTGTCTGCTGCTGGCTGGCTGG 3780 | 4801 | TGATAGGCAATTAAACTCTTAAAGAAATTATGTTAGATGAACTTCAAACT 4860 |
| Db | 3721 | TATACTATGATAAAGTTGTTATGCTGGAAAGTGTCTGCTGCTGGCTGGCTGG 3780 | 4801 | TGATAGGCAATTAAACTCTTAAAGAAATTATGTTAGATGAACTTCAAACT 4860 |
| Qy | 3781 | TAATGTGTCGCCCTAGTCGCTGAGTATGCTTAACTGTTAGCTAGCTTGTAA 3840 | 4861 | GCTGTCCTTCTTGTGTTTGTGTTTGTGTTTGTGTTTGTGTTTGTGTTTGTG 4920 |
| Db | 3781 | TAATGTGTCGCCCTAGTCGCTGAGTATGCTTAACTGTTAGCTAGCTTGTAA 3840 | 4861 | GCTGTCCTTCTTGTGTTTGTGTTTGTGTTTGTGTTTGTGTTTGTGTTTGTG 4920 |
| Qy | 3841 | TTTAAAAAAATTGTTAAATTSCATGCTGGAACTTCTGCTGCTGCTGGCTGG 3900 | 4921 | TAGGACAGTCATGCTTGTGAAATTCTGCTGAGGATATTGTGTTTGTG 4980 |
| Db | 3841 | TTTAAAAAAATTGTTAAATTSCATGCTGGAACTTCTGCTGCTGCTGGCTGG 3900 | 4921 | TAGGACAGTCATGCTTGTGAAATTCTGCTGAGGATATTGTGTTTGTG 4980 |
| Qy | 3901 | AGATGAAATTCACTTACCTTACCTCTGCTGGCTGGCTGGCTGGCTGGCTGG 3960 | 5041 | ATAATAACAAATACCCGCTATGCTGTAATTCTGCAATGTCCTCTGTGAC 5040 |
| Db | 3901 | AGATGAAATTCACTTACCTCTGCTGGCTGGCTGGCTGGCTGGCTGGCTGG 3960 | 5041 | ATAATAACAAATACCCGCTATGCTGTAATTCTGCAATGTCCTCTGTGAC 5040 |
| Qy | 3961 | TCCATTATTCTGCTTTCATTTTCACTTCACTTCACTTCACTTCACTTCACT 4020 | 5101 | GGAAATACCAATAAGTAACTACCAAAAGTTGACTACTAAAGGGATATGCT 5160 |
| Db | 3961 | TCCATTATTCTGCTTTCATTTTCACTTCACTTCACTTCACTTCACTTCACT 4020 | 5101 | GGAAATACCAATAAGTAACTACCAAAAGTTGACTACTAAAGGGATATGCT 5160 |
| Qy | 4081 | ACAAATATTGAGCATATTCCATTTCATTAAATCTCCACAAAGTTATTGT 4140 | 5161 | GAAAAGCTATAAAAGAAUUTTCAGCATGATTTCATATTGTGCTTCCACCT 5220 |
| Db | 4081 | ACAAATATTGAGCATATTCCATTTCATTAAATCTCCACAAAGTTATTGT 4140 | 5161 | GAAAAGCTATAAAAGAAUUTTCAGCATGATTTCATATTGTGCTTCCACCT 5220 |

| | | | | | | | | |
|----|------|---|------|--|----|------|--|------|
| Db | 1681 | AGAGTGAAAGGGGCCGCTGATTAATACACTAAGTCATAAGCCAGGACT | 1740 | | Qy | 2821 | CAGAGGAAAGTCCAAAAGGGACCTGAACTGTGAAACTGATGAAAAGTTGTTACTCTGGGTAGTC | 2880 |
| Qy | 1741 | GTTTGGTAAACTGGTCACTTATCCTTAACCTAAACTAACATGACTAATT | 1800 | | Db | 2821 | CAGAGGAAAGTCCAAAAGGGACCTGAACTGTGAAACTGATGAAAAGTTGTTACTCTGGGTAGTC | 2880 |
| Db | 1741 | GTTTGGTAAACTGGTCACTTATCCTTAACCTAAACTAACATGACTAATT | 1800 | | Qy | 2881 | GCATGACATCTTGACTTATCCTTAACCTAAACTAACATGACTAATT | 2940 |
| Qy | 1801 | AGTTACTAAGTCCTTGACTTATCCTTAACCTAACTGGCTTACCTTAT | 1860 | | Db | 2881 | GCATGACATCTTGACTTATCCTTAACCTAAACTAACATGACTAATT | 2940 |
| Db | 1801 | AGTTACTAAGTCCTTGACTTATCCTTAACCTAACTGGCTTACCTTAT | 1860 | | Qy | 2941 | TTATTGTGAGTTGCTGTTGTTATTATTTATATGAAATAGTCATAAT | 3000 |
| Qy | 1861 | TGACAGTATTATTGGAAAACCTCTTAACCTCATAGTCCTTATCCCTT | 1920 | | Db | 2941 | TTATTGTGAGTTGCTGTTGTTATTATTTATATGAAATAGTCATAAT | 3000 |
| Db | 1861 | TGACAGTATTATTGGAAAACCTCTTAACCTCATAGTCCTTATCCCTT | 1920 | | Qy | 3001 | TTACTAAAGGCCPACCATGGCGCATTTTACATTGTCCTCPAACTCTTGATG | 3060 |
| Qy | 1921 | TGAAACAAAGAGAGCTCAAACTACATAATGATTATTAGTCCTTGTGT | 1980 | | Db | 3001 | TTACTAAAGGCCPACCATGGCGCATTTTACATTGTCCTCPAACTCTTGATG | 3060 |
| Db | 1921 | TGAAACAAAGAGAGCTCAAACTACATAATGATTATTAGTCCTTGTGT | 1980 | | Qy | 3061 | AGATGATCAGATGGATACCTGGCTTGAAGTGAATATGATGTTACATCTATATC | 3120 |
| Qy | 1981 | CTATAATAGTCCTCCAGAGGAGCTTAAACTCTCCATTAAAGTCCTTGTG | 2040 | | Db | 3061 | AGATGATCAGATGGATACCTGGCTTGAAGTGAATATGATGTTACATCTATATC | 3120 |
| Db | 1981 | CTATAATAGTCCTCCAGAGGAGCTTAAACTCTCCATTAAAGTCCTTGTG | 2040 | | Qy | 3121 | TATATCTATATCPTATATCPTATATCPTATATCPTATATCPTATATC | 3180 |
| Qy | 2041 | TGCCAACTTGGCAGAAATTCCAAATATCTGATTTCTGCTTGTG | 2100 | | Db | 3121 | TATATCTATATCPTATATCPTATATCPTATATCPTATATCPTATATC | 3180 |
| Db | 2041 | TGCCAACTTGGCAGAAATTCCAAATATCTGATTTCTGCTTGTG | 2100 | | Qy | 3181 | AAATATGTGTTGTAAGTTAAGATTAAAGTAAAGTAAAGTAAAGTAA | 3240 |
| Qy | 2101 | TTCAAACACTGATTTCTCTTACCTGGTCACTTGTGCTTGTG | 2160 | | Db | 3181 | AAATATGTGTTGTAAGTTAAGATTAAAGTAAAGTAAAGTAAAGTAA | 3240 |
| Db | 2101 | TTCAAACACTGATTTCTCTTACCTGGTCACTTGTGCTTGTG | 2160 | | Qy | 3241 | GTAAACCCCTTCTCTGCTGCTGCTGAAATAACATATTATTCATTCATC | 3300 |
| Qy | 2161 | TTTCCACACCCTTACATTTCCTGGAAACCTACATGTTGCTTGTG | 2220 | | Db | 3241 | GTAAACCCCTTCTCTGCTGCTGCTGAAATAACATATTATTCATTCATC | 3300 |
| Db | 2161 | TTTCCACACCCTTACATTTCCTGGAAACCTACATGTTGCTTGTG | 2220 | | Qy | 3301 | AAATGTGTTGCTGAAAGCAGCTGTGAAAGCCTTGGGATATTGCGC | 3360 |
| Qy | 2221 | TCTTAATTTCCTTCAGATCTTCATGAGTGTCTTCTGCTG | 2280 | | Db | 3301 | AAATGTGTTGCTGAACTGCTGCTGCTGAAAGCCTTGGGATATTGCGC | 3360 |
| Db | 2221 | TCTTAATTTCCTTCAGATCTTCAGATCTTCAGATCTTCAG | 2280 | | Qy | 3361 | CATGCTGGCAACGCTTATAGTTGCTTCAAAACTCTTCACTTAATCT | 3420 |
| Qy | 2281 | CCTCCCAAAGATGTCATGAGTCTCCCTTTTATCTACTAATCACCA | 2340 | | Db | 3361 | CATGCTGGCAACGCTTATAGTTGCTTCAAAACTCTTCACTTAATCT | 3420 |
| Db | 2281 | CCTCCCAAAGATGTCATGAGTCTCCCTTTTATCTACTAATCACCA | 2340 | | Qy | 3421 | AATACCTCATGACTATTGCTTTCAGGTTTCACTACAAATAACTCTGCTTTCATA | 3480 |
| Qy | 2341 | TGTTGTGATTAATGACTAATGCTGTTCTGTTGATTAATGCTG | 2400 | | Db | 3421 | AATACCTCATGACTATTGCTTTCAGGTTTCACTACAAATAACTCTGCTTTCATA | 3480 |
| Db | 2341 | TGTTGTGATTAATGACTAATGCTGTTCTGTTGATTAATGCTG | 2400 | | Qy | 3481 | TATTTGAGTAAGTCCTCCCTGGAAAGTAGTAAAGTAAAGTAA | 3540 |
| Qy | 2401 | TACATGGTCTCTGTTGCTGTTGATTAATGCTGAAATGCA | 2460 | | Db | 3481 | TATTTGAGTAAGTCCTCCCTGGAAAGTAGTAAAGTAAAGTAA | 3540 |
| Db | 2401 | TACATGGTCTCTGTTGCTGTTGATTAATGCTGAAATGCA | 2460 | | Qy | 3541 | TGGATCCAACCGTATAGCAAGCTTGTGCTGCTGAACTGCAAG | 3600 |
| Qy | 2461 | GACTAACATTTAATGAAATGAAATGAAACCCATCTATCGAGTGTG | 2520 | | Db | 3541 | TGGATCCAACCGTATAGCAAGCTTGTGCTGCTGAACTGCAAG | 3600 |
| Db | 2461 | GACTAACATTTAATGAAATGAAATGAAACCCATCTATCGAGTGTG | 2520 | | Qy | 3601 | GCTCTGTTGTTGTTAACATGTTCTCTGGGACAAACCTAGGAGAACGCTAGCAGAA | 3660 |
| Qy | 2521 | AGACCGGTTCTGAGCCATTATTTGTTGAAATGAAACCTGAACT | 2580 | | Db | 3601 | GCTCTGTTGTTGTTAACATGTTCTCTGGGACAAACCTAGGAGAACGCTAGCAGAA | 3660 |
| Db | 2521 | AGACCGGTTCTGAGCCATTATTTGTTGAAATGAAACCTGAACT | 2580 | | Qy | 3661 | AATGGACAAAACCTGAACTGTTGCTGAACTGCTGAACTGCTGAA | 3720 |
| Qy | 2581 | GAGGTACTATCACTATCCTTATGTTGAAAGATAAGCCAGAAATGAA | 2640 | | Db | 3661 | AATGGACAAAACCTGAACTGCTGAACTGCTGAACTGCTGAA | 3720 |
| Db | 2581 | GAGGTACTATCACTATCCTTATGTTGAAAGATAAGCCAGAAATGAA | 2640 | | Qy | 3721 | TATCTATGTAATATGTTGTTGCTGAACTGCTGAACTGCTGAA | 3780 |
| Qy | 2641 | AACTCACCAAAAGTCATGCTGAACTGCTGAACTGCTGAACTGCT | 2700 | | Db | 3721 | TATCTATGTAATATGTTGTTGCTGAACTGCTGAACTGCTGAA | 3780 |
| Db | 2641 | AACTCACCAAAAGTCATGCTGAACTGCTGAACTGCTGAACTGCT | 2700 | | Qy | 3781 | TAATGTGTCCTAGTGAATGTTAAATGACTGACTGACTGACTG | 3840 |
| Qy | 2701 | TTACATGTTGCTCTGCTGCTGCTGCTGCTGCTGCTGCTGCTG | 2760 | | Db | 3781 | TAATGTGTCCTAGTGAATGTTAAATGACTGACTGACTGACTG | 3840 |
| Db | 2701 | TTACATGTTGCTCTGCTGCTGCTGCTGCTGCTGCTGCTGCTG | 2760 | | Qy | 3841 | TTTAAATTTAAATTTGTTAAATGCTAGTGCCTTCATGAGAG | 3900 |
| Qy | 2761 | CTGCTCGTAAAGGGCAATATGGGAGGAACTGGAGGAACTGGAG | 2820 | | Db | 3841 | TTTAAATTTAAATTTGTTAAATGCTAGTGCCTTCATGAGAG | 3900 |
| Db | 2761 | CTGCTCGTAAAGGGCAATATGGGAGGAACTGGAGGAACTGGAG | 2820 | | | | | |

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|------|--|------|----|--|---|------|
| 3901 | AGATAGAATTAACTTCACTTACTTACCACTTAGAACCCATTAAACTTGTG | 3960 | Db | 4981 | AAATAAAATAAAGTTGCACTTTCCTGCCCCAAAGAGCTCTGTGTCCTGAC | 504 |
| 3901 | AGATAGAATTAACTTCACTTACTTACCACTTAGAACCCATTAAACTTGTG | 3960 | Db | 5041 | ATAAAATAACAAACCGPATGCTTTAATTATGGCAATTGCCCCATTTCAACCTAA | 510 |
| 3961 | TCCATTATTCGCTTTATTCAACATTTCAGGAACTTACAGGGAG | 4020 | Db | 5041 | ATAAAATAACAAACCGPATGCTTTAATTATGGCAATTGCCCCATTTCAACCTAA | 510 |
| 3961 | TCCATTATTCGCTTTATTCAACATTTCAGGAACTTACAGGGAG | 4020 | Qy | 5101 | GGAATTAACCTAAAGTAAACAGATAACCAAAAGTTACTAGTTAACGGGCATTSCT | 516 |
| 4021 | GTACATGATAACAAATGAGAGACTCCCATGTTCTGTTTCAGTTA | 4080 | Db | 5101 | GGAATTAACCTAAAGTAAACAGATAACCAAAAGTTACTAGTTAACGGGCATTSCT | 516 |
| 4021 | GTACATGATAACAAATGAGAGACTCCCATGTTCTGTTTCAGTTA | 4080 | Qy | 5161 | GAAGAGTAAAGAATTTCAGATGTTCCATATGCTATATGCTTACACACTGCCAT | 522 |
| 4081 | ACAATATATGAGCATATTCCATTCAATTAACTTCACAAGTTATTGTAT | 4140 | Db | 5161 | GAAGAGTAAAGAATTTCAGATGTTCCATATGCTTACACACTGCCAT | 522 |
| 4081 | ACAATATATGAGCATATTCCATTCAATTAACTTCACAAGTTATTGTAT | 4140 | Qy | 5221 | AACA | 5224 |
| 4141 | GGCTGTTATACCCCTACTTATGATGTTACCATTAATTTCCTGGGTAA | 4200 | Db | 5221 | ANCA | 5224 |
| 4141 | GGCTGTTATACCCCTACTTATGATGTTACCATTAATTTCCTGGGTAA | 4200 | Qy | RESULT 7 | | |
| 4201 | TTTGATTTTATAATCTTACCCCTAGAATAATGAAACCCCTGAAAGCTT | 4260 | Db | US-10-139-089-5 | | |
| 4201 | TTTGATTTTATAATCTTACCCCTAGAATAATGAAACCCCTGAAAGCTT | 4260 | Qy | / Sequence 5 , Application US/10139089 | | |
| 4261 | GGTGCCTGGTCTCACTCCACAGATTCTGTTAACGCTAGGCA | 4320 | Db | / Publication No. US20030152553A1 | | |
| 4261 | GGTGCCTGGTCTCACTCCACAGATTCTGTTAACGCTAGGCA | 4320 | Qy | / GENERAL INFORMATION: | | |
| 4321 | TTGGAAATTCAAAGTTCCCAGTGTACCTGAAAGTCGGAAACCCTTG | 4380 | Db | / APPLICANT: Henderson, D. R. | | |
| 4321 | TTGGAAATTCAAAGTTCCCAGTGTACCTGAAAGTCGGAAACCCTTG | 4380 | Qy | / TITLE OF INVENTION: TISSUE SPECIFIC VIRAL VECTORS | | |
| 4381 | TAGACAGGGATGATAGGGTGGCCTTACATCATTAGPATTACATCAT | 4440 | Db | / FILE REFERENCE: 348020000221 | | |
| 4381 | TAGACAGGGATGATAGGGTGGCCTTACATCATTAGPATTACATCAT | 4440 | Qy | / CURRENT APPLICATION NUMBER: US/10/139, 089 | | |
| 4441 | CATCTTGAGTTGCTTAAGTGTGCTTAACTGCTTAACTGCTAA | 4500 | Db | / CURRENT FILING DATE: 2003-05-02 | | |
| 4441 | CATCTTGAGTTGCTTAAGTGTGCTTAACTGCTTAACTGCTAA | 4500 | Qy | / PRIORITY APPLICATION NUMBER: 08/669, 753 | | |
| 4501 | TAAAATTATATAGGACTTCTGGTTATTAGGGCTTGTGCTTAAGGCC | 4560 | Db | / PRIORITY FILING DATE: 1996-06-26 | | |
| 4501 | TAAAATTATATAGGACTTCTGGTTATTAGGGCTTGTGCTTAAGGCC | 4560 | Qy | / PRIORITY APPLICATION NUMBER: 08/495, 034 | | |
| 4561 | ATACATCGATACTAAATCTTAAATGCTACATTTACATTTGAAAG | 4620 | Db | / PRIORITY FILING DATE: 1995-06-27 | | |
| 4561 | ATACATCGATACTAAATCTTAAATGCTACATTTGAAAG | 4620 | Qy | / PRIORITY APPLICATION NUMBER: 09/509, 591 | | |
| 4621 | TGTTCTGAGCTAAACATGACGCTTAAATGCTACATTTGAAAGT | 4680 | Db | / PRIORITY FILING DATE: 2000-06-02 | | |
| 4621 | TGTTCTGAGCTAAACATGACGCTTAAATGCTACATTTGAAAGT | 4680 | Qy | / PRIORITY APPLICATION NUMBER: 09/151, 376 | | |
| 4681 | ATTATTCGCAACTTGGGACAACCTCATCTCTGAAATTCTGAGTATT | 4740 | Db | / PRIORITY FILING DATE: 1998-04-10 | | |
| 4681 | ATTATTCGCAACTTGGGACAACCTCATCTCTGAAATTCTGAGTATT | 4740 | Qy | / PRIORITY APPLICATION NUMBER: 09/033, 428 | | |
| 4741 | GTATATTGCAAGATGAAAGTGTGAACTTGTGCTGCGCATA | 4800 | Db | / PRIORITY FILING DATE: 1998-01-02 | | |
| 4741 | GTATATTGCAAGATGAAAGTGTGAACTTGTGCTGCGCATA | 4800 | Qy | / PRIORITY APPLICATION NUMBER: 60/039, 597 | | |
| 4801 | TGATAGGCAATTAACTGTTAAAGAATTAAATGCTATTGAAATTCT | 4860 | Db | / PRIORITY FILING DATE: 1997-03-03 | | |
| 4801 | TGATAGGCAATTAACTGTTAAAGAATTAAATGCTATTGAAATTCT | 4860 | Qy | / PRIORITY APPLICATION NUMBER: 09/033, 555 | | |
| 4861 | GGCTCTTCTTCTTATGGCTTCAATTAACTTAATTCTGCACT | 4920 | Db | / PRIORITY FILING DATE: 1998-04-02 | | |
| 4861 | GGCTCTTCTTATGGCTTCAATTAACTTAATTCTGCACT | 4920 | Qy | / PRIORITY APPLICATION NUMBER: 60/039, 763 | | |
| 4921 | TAGGACAGTCATGCTTCAATTAACTTAATTCTGCACT | 4980 | Db | / PRIORITY FILING DATE: 1997-03-03 | | |
| 4921 | TAGGACAGTCATGCTTCAATTCTGAGGAAATTGTGTTGAGTAT | 4980 | Qy | / NUMBER OF SEQ ID NOS: 71 | | |
| 4981 | AAATAATAATAAGTTGCAAGTAACTTAAACTTAATTCTGCACT | 5040 | Db | / SOFTWARE: Ptent.In Ver. 2.0 | | |
| 4981 | AAATAATAATAAGTTGCAAGTAACTTAAACTTAATTCTGCACT | 5040 | Qy | / SEQ ID NO: 5 | | |
| 4981 | AAATAATAATAAGTTGCAAGTAACTTAAACTTAATTCTGCACT | 5040 | Qy | / LENGTH: 5224 | | |
| 4981 | AAATAATAATAAGTTGCAAGTAACTTAAACTTAATTCTGCACT | 5040 | Qy | / TYPE: DNA | | |
| 4981 | AAATAATAATAAGTTGCAAGTAACTTAAACTTAATTCTGCACT | 5040 | Qy | / ORGANISM: Homo sapiens | | |
| 4981 | AAATAATAATAAGTTGCAAGTAACTTAAACTTAATTCTGCACT | 5040 | Qy | / US-10-139-089-5 | | |
| 4981 | AAATAATAATAAGTTGCAAGTAACTTAAACTTAATTCTGCACT | 5040 | Qy | Query Match 100.0% Score 5224; DB 16; Length 5224; | | |
| 4981 | AAATAATAATAAGTTGCAAGTAACTTAAACTTAATTCTGCACT | 5040 | Qy | Best Local Similarity 100.0% Pred. No. 0; | | |
| 4981 | AAATAATAATAAGTTGCAAGTAACTTAAACTTAATTCTGCACT | 5040 | Qy | Matches 5224; Conservative 0; Mismatches 0; Indels 0; Gaps 0 | | |
| 4981 | AAATAATAATAAGTTGCAAGTAACTTAAACTTAATTCTGCACT | 5040 | Qy | 1 GAATTCCTAGAAATAAGGGGTAGGGCTGGTAAATTCTGTTTACCCCCATTAGGT | 60 | |
| 4981 | AAATAATAATAAGTTGCAAGTAACTTAAACTTAATTCTGCACT | 5040 | Qy | 1 GAATTCCTAGAAATAAGGGGTAGGGCTGGTAAATTCTGTTTACCCCCATTAGGT | 60 | |
| 4981 | AAATAATAATAAGTTGCAAGTAACTTAAACTTAATTCTGCACT | 5040 | Qy | 61 GAGATAGGATGGGTTAAATGTTGCTTCAACACATOCATTCTAGAATAAGG 120 | | |
| 4981 | AAATAATAATAAGTTGCAAGTAACTTAAACTTAATTCTGCACT | 5040 | Qy | 61 GAGATAGGATGGGTTAAATGTTGCTTCAACACATOCATTCTAGAATAAGG 120 | | |
| 4981 | AAATAATAATAAGTTGCAAGTAACTTAAACTTAATTCTGCACT | 5040 | Qy | 121 AACAGACTATGGGTTGGAGCACTTGGTTCTCATACACTTCATGTTTCT | 180 | |

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|------|--|------|----------------|-------|--|------|
| 121 | AACAGACTATGGGTGGAGGACTTTGGATTGTCCTCATAAACACTTGGGTGTCATCP | 180 | Qy | 1.261 | ACTTAGGACTAAACTTGACCAATAAACATGAGATGAGGATTCAGATGAGATGTCAT | 1320 |
| 181 | GTTCTATGGGTGGTTTAAGCTTGGCAACTTGGCAACTTGGCTCC | 240 | D _b | 1.261 | ACTTAGGACTAAACTTGACCAATAAACATGAGATGAGGATTCAGATGAGATGTCAT | 1320 |
| 181 | GTTCTATGGGTGGTTTAAGCTTGGCAACTTGGCTCC | 240 | Qy | 1.321 | ACAACACTTCGGTCAAAGCTCCTTATTGCTGTTCTGAGATGAGATGAGATGTCAT | 1380 |
| 241 | AAGCCCCAAGGACTCTGCTCTTTCATATCTGTTGGGCCCTCTGGGCTTGAATATCT | 300 | D _b | 1.321 | ACAACACTTCGGTCAAAGCTCCTTATTGCTGTTCTGAGATGAGATGAGATGTCAT | 1380 |
| 241 | AAGCCCCAAGGACTCTGCTCTTTCATATCTGTTGGCAACTTGGCTCC | 300 | Qy | 1.381 | GGTTCTCTTTCACTGTPATTTTCTCACACCACCTCACATGGCTAACATAACTGGTC | 1440 |
| 301 | GAGAAAATATAAACATTCATAATGGTCTGGTCAAGTAGTGTAGGATGTCAT | 360 | D _b | 1.381 | GGTTCTCTTTCACTGTPATTTTCTCACACCACCTCACATGGCTAACATAACTGGTC | 1440 |
| 301 | GAGAAAATATAAACATTCATAATGGTCTGGTCAAGTAGTGTAGGATGTCAT | 360 | Qy | 1.441 | TGCCAGCTPATGATTCCTCACATCTAGCTGTTCTAGAAAGATAAA | 1500 |
| 361 | TCTTGTGATCAATGATGATGAACTATTAGTGTATAAACTCTTAGACAACTT | 420 | D _b | 1.441 | TGCCAGCTPATGATTCCTCACATCTAGCTGTTCTAGAAAGATAAA | 1500 |
| 361 | TCTTGTGATCAATGATGATGAACTATTAGTGTATAAACTCTTAGACAACTT | 420 | Qy | 1.501 | AGTACTTAAATGCACATCAAAGCTGTTCACTTGGGGCTTAAAGAGCTTCAACATA | 1560 |
| 421 | GGGGTAGGGGGTGTACTATTCATAATTAAAGGACTATTCTTATTGTTA | 480 | D _b | 1.501 | AGTACTTAAATGCACATCAAAGCTGTTCACTTGGGGCTTAAAGAGCTTCAACATA | 1560 |
| 421 | GGGGTAGGGGGTGTACTATTCATAATTAAAGGACTATTCTTATTGTTA | 480 | Qy | 1.561 | CAAACCGGGAGTTTGCCTGGTAAATGTTCTAAATGTTCTGACATAGGTGCC | 1620 |
| 481 | TGCTTGTGACAAATGTTTTCGGGACCAAGGATCACAAAGTAGCTTCATAIT | 540 | D _b | 1.561 | CAAACCGGGAGTTTGCCTGGTAAATGTTCTAAATGTTCTGACATAGGTGCC | 1620 |
| 481 | TGCTTGTGACAAATGTTTTCGGGACCAAGGATCACAAAGTAGCTTCATAIT | 540 | Qy | 1.621 | TCTTGTCTCTAAATCTATTACTTTAGGCCCTGTCTCATCCCACTATGGGGAGATG | 1680 |
| 541 | AAGAAGTTAAATGGTCCAGGATAATTACATAGCTTACAATGACTGATAACATA | 600 | D _b | 1.621 | TCTTGTCTCTAAATCTATTACTTTAGGCCCTGTCTCATCCCACTATGGGGAGATG | 1680 |
| 541 | AAGAAGTTAAATGGTCCAGGATAATTACATAGCTTACAATGACTGATAACATA | 600 | Qy | 1.681 | AGAGTGAAGGGGCGCTTAAATTAACCTAACTTAACTGAACTAGGCAACT | 1740 |
| 601 | ACAAGAGGTTCATGAGAAATAATCTGAAGGTTAAATAGTGTCAAGGTGAGAGC | 660 | D _b | 1.681 | AGAGTGAAGGGGCGCTTAAATTAACCTAACTTAACTGAACTAGGCAACT | 1740 |
| 601 | ACAAGAGGTTCATGAGAAATAATCTGAAGGTTAAATAGTGTCAAGGTGAGAGC | 660 | Qy | 1.741 | GTGTTGGTAAACCTGTCACCTTAACTTAACTGAACTATCCAAAACCTGAACTT | 1800 |
| 661 | CTCTTCTCTAGCTAGACTATCACAAATCACATTCACTGATTAATTTGTTA | 720 | D _b | 1.741 | GTGTTGGTAAACCTGTCACCTTAACTTAACTGAACTATCCAAAACCTGAACTT | 1800 |
| 661 | CTCTTCTCTAGCTAGACTATCACAAATCACATTCACTGATTAATTTGTTA | 720 | Qy | 1.801 | AGTTACTAAGTCTTGTGACTTTATCTCATCCATCAGCTTATCCGCTT | 1860 |
| 721 | TAAGGGTTGGTACATTGGTCAAGCTTCACTGAGAAAGCTGAAATTGAAAC | 780 | D _b | 1.801 | AGTTACTAAGTCTTGTGACTTTATCTCATCCATCAGCTTATCCGCTT | 1860 |
| 721 | TAAGGGTTGGTACATTGGTCAAGCTTCACTGAGAAAGCTGAAATTGAAAC | 780 | Qy | 1.861 | TTGACAGTTATTGCGGAAACTCTCTTACTGTTCTCTTATCCGCTT | 1920 |
| 781 | ATTTCTCACTAACCAACCACTTCAACAAACAAAAATGAGAAATCTGAGAAAC | 840 | D _b | 1.861 | TTGACAGTTATTGCGGAAACTCTCTTACTGTTCTCTTATCCGCTT | 1920 |
| 781 | ATTTCTCACTAACCAACCACTTCAACAAACAAAAATGAGAAATCTGAGAAAC | 840 | Qy | 1.921 | TGAAACAAAAGAGACAGTTCAAAATCACAAATATGATTTTTATTGCTCTT | 1980 |
| 841 | CTGAGATAAGGAAAGGAAATTCTCAACCCACAGTAACTGCTCTGAGAA | 900 | D _b | 1.921 | TGAAACAAAAGAGACAGTTCAAAATCACAAATATGATTTTTATTGCTCTT | 1980 |
| 841 | CTGAGATAAGGAAAGGAAATTCTCAACCCACAGTAACTGCTCTGAGAA | 900 | Qy | 2.041 | TGCCAAACTGCAATTTCCTCTCCTAACGTTCTGAGATGCGCC | 2100 |
| 901 | TTCATATGCTATAATTAACTAACTCTGAGATATTTCATTATATCAAAATA | 960 | D _b | 2.041 | TGCCAAACTGCAATTTCCTCTCCTAACGTTCTGAGATGCGCC | 2100 |
| 901 | TTCATATGCTATAATTAACTAACTCTGAGATATTTCATTATATCAAAATA | 960 | Qy | 2.101 | TTCCAAAACCTGCAATTTCCTCTCCTAACGTTCTGAGATGCGCC | 2100 |
| 1021 | ACTTGAGGGGATAGATCATTTCTCATGAGAAATTAAACGATTGAAATT | 1080 | D _b | 2.101 | TTCCAAAACCTGCAATTTCCTCTCCTAACGTTCTGAGATGCGCC | 2100 |
| 1021 | ACTTGAGGGGATAGATCATTTCTCATGAGAAATTAAACGATTGAAATT | 1080 | Qy | 2.161 | TTTCACCCACCTTTCATCTGGACACTATACCTCCCTCTCATTTGCCAC | 2220 |
| 1081 | TTGCCCTGTCATACAGTCAATTGACCTTCACTGAGATTTAGTTGAAAT | 1140 | D _b | 2.161 | TTTCACCCACCTTTCATCTGGACACTATACCTCCCTCTCATTTGCCAC | 2220 |
| 1081 | TTGCCCTGTCATACAGTCAATTGACCTTCACTGAGATTTAGTTGAAAT | 1140 | Qy | 2.221 | CTTCAAAAGATGTCATGAGCTCTTCATCTAACGATTCATCACACCA | 2340 |
| 1141 | CTTCTCTATAACCAAAAGTCACTTCACTGAGAAATTAAACGATTGAAATT | 1200 | D _b | 2.221 | CTTCAAAAGATGTCATGAGCTCTTCATCTAACGATTCATCACACCA | 2340 |
| 1141 | CTTCTCTATAACCAAAAGTCACTTCACTGAGAAATTAAACGATTGAAATT | 1200 | Qy | 2.281 | CCTCCAAGATGTCATGAGCTCTTCATCTAACGATTCATCACACCA | 2340 |
| 1201 | TATGAAAAGTAAACGGAACTGAGAACTTACATCAATGCAAAAGCTGACTCT | 1260 | D _b | 2.281 | CCTCCAAGATGTCATGAGCTCTTCATCTAACGATTCATCACACCA | 2340 |

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|----|------|---|------|----|------|---|------|
| Db | 721 | TAAGGGTTGGTACATTGTCAGGATTGTGGAGAAGGAGTCATAATTTGAAAAC | 780 | Qy | 1861 | TTGACAGTATTGCGAAACTCCCTAACCTGCTCCTTATCATAGCTTATCCCTTT | 1920 |
| Qy | 781 | ATTTCAACTAACCAAACCCAAATCTAACACAAAAAAATGAAATCTCAGAAAACA | 840 | Db | 1861 | TGACAGTATTGCGAAACTCCCTAACCTGCTCCTTATCATAGCTTATCCCTTT | 1920 |
| Db | 781 | ATTTCAACTAACCAAACCCAAATCTAACACAAAAAAATGAAATCTCAGAAAACA | 840 | Qy | 1921 | TGAACAAAGAGACAGTTCAAAATAAAATAATGATTTTTATTAGCTCCCTTGTGT | 1980 |
| Qy | 841 | GTGAGATAAGAAGGAAATTCTTCACAACCCACGTATAGCTCAACTGCTTGAGAA | 900 | Db | 1921 | TGAACAAAGAGACAGTTCAAAATAAAATAATGATTTTTATTAGCTCCCTTGTGT | 1980 |
| Db | 841 | GTGAGATAAGAAGGAAATTCTTCACAACCCACGTATAGCTCAACTGCTTGAGAA | 900 | Qy | 1981 | CTATATAAGTCCAGAGGAGTTAAACCTCAATTAAAGTCTTGAAGTGTGCCT | 2040 |
| Qy | 901 | GTATATCTTAAATTAACTAACTGTAATAATTGTCATT | 960 | Db | 1981 | CTATATAAGTCCAGAGGAGTTAAACCTCAATTAAAGTCTTGAAGTGTGCCT | 2040 |
| Qy | 901 | GTATATCTTAAATTAACTAACTGTAATAATTGTCATT | 960 | Qy | 2041 | TGCCAACTTGCAGGAAATTCCCAATAATCPAGTATTTCPACTATAAACCTTGTGCCT | 2100 |
| Db | 961 | TTAATGCTTAAAGTACACTAACATCAGCTAATAATTCAAAATAA | 1020 | Db | 2041 | TGCCAACTTGCAGGAAATTCCCAATAATCPAGTATTTCPACTATAAACCTTGTGCCT | 2100 |
| Db | 961 | TTAATGCTTAAAGTACAGGCAATTAGATATTATCAAAATAA | 1020 | Qy | 2101 | TTCAAAAACTGCATTCTCATTCCCTAACCTGCTTACAGTGTGCCTTACGGTRGGT | 2160 |
| Qy | 1021 | ACTTGAGGGTGGATAGATATTCTCATATATGAGAAAATTAAACAGTTGAAATT | 1080 | Db | 2101 | TTCAAAAACTGCATTCTCATTCCCTAACCTGCTTACAGTGTGCCTTACGGTRGGT | 2160 |
| Db | 1021 | ACTTGAGGGTGGATAGATATTCTCATATATGAGAAAATTAAACAGTTGAAATT | 1080 | Db | 2161 | TTTCACCAACCTTTCATTTTCAATTCCGAAACATTAACCCCTTCATTTGACC | 2220 |
| Qy | 1081 | TTGCGCTGTCAGTTACTGTTCCATGTTCTGACTGCTTCACAGCT | 1140 | Db | 2161 | TTTCACCAACCTTTCATTTTCAATTCCGAAACATTAACCCCTTCATTTGACC | 2220 |
| Db | 1081 | TTGCGCTGTCAGTTACTGTTCCATGAGCTTCACTGAGTT | 1140 | Qy | 2221 | TCTAATTTCCTTCAGATCTCATAGATTTCACTTCAAGAGCTTATCTGACC | 2280 |
| Qy | 1141 | CTTCTPAATAACCAAACTTCAGTTACTGTTCCATGTTCTGACTGCTTCACAGCT | 1200 | Db | 2221 | TCTAATTTCCTTCAGATCTCATAGATTTCACTTCAAGAGCTTATCTGACC | 2280 |
| Db | 1141 | CTTCTPAATAACCAAACTTCAGTTACTGTTCCATGAGCTTCACTGAGTT | 1200 | Qy | 2281 | CCTCCAAGAAGTGTCTGACTGCTTCCTTCACTTCAAGAGCTTATCTGACC | 2340 |
| Qy | 1201 | TATGAAAAGTAAACGSAATCAGAAATTACATCAATCAGCAAGCATGCTGAACTCTGT | 1260 | Db | 2281 | CCTCCAAGAAGTGTCTGACTGCTTCCTTCACTTCAAGAGCTTATCTGACC | 2340 |
| Db | 1201 | TATGAAAAGTAAACGSAATCAGAAATTACATCAATCAGCAAGCATGCTGAACTCTGT | 1260 | Qy | 2341 | TGTTGTGATPACTGATACTATTGCTGTTCTCTGTTCTGTTCTGTTCTGTTCTG | 2400 |
| Qy | 1261 | ACTTAGGACTAACTTGTGAGGATAACACATAGATTGAGGATTTGTGTGTTGAT | 1320 | Db | 2341 | TGTTGTGATPACTGATACTATTGCTGTTCTCTGTTCTGTTCTGTTCTGTTCTG | 2400 |
| Db | 1261 | ACTTAGGACTAACTTGTGAGGATAACACATAGATTGAGGATTTGTGTGTTGAT | 1320 | Qy | 2401 | TACATGGCTGCTGTCCTGTTGCTGTTGCTGTTGCTGTTGCTGTTGCTG | 2460 |
| Qy | 1321 | ACAAACTCTGTCTTCAAGCTCTCTTCAAGCTTCAATCTGTTGAGGATCTTGTCTCAT | 1380 | Db | 2401 | TACATGGCTGCTGTCCTGTTGCTGTTGCTGTTGCTGTTGCTGTTGCTG | 2460 |
| Db | 1321 | ACAAACTCTGTCTTCAAGCTCTCTTCAAGCTTCAATCTGTTGAGGATCTTGTCTCAT | 1380 | Qy | 2461 | GACTAACATTTTATGAAATAATAAAACCCCATCTATGAGCTGCTACTTTGTGCA | 2520 |
| Qy | 1381 | GTTTCTCTTCTACGCTATCTTCTCAACCACTCACATGCTCAATACTGTC | 1440 | Db | 2461 | GACTAACATTTTATGAAATAATAAAACCCCATCTATGAGCTGCTACTTTGTGCA | 2520 |
| Db | 1381 | GTTTCTCTTCTACGCTATCTTCTCAACCACTCACATGCTCAATACTGTC | 1440 | Qy | 2521 | AGACCGGTTCTGAGSCATTATATTGTTATTCTCATTTAACCTGAAAG | 2580 |
| Qy | 1441 | TGCAAGCTTATGTTCCAAATATCTATCTCTAGCTCAATTGTTCCAGAGTAA | 1500 | Db | 2521 | AGACCGGTTCTGAGSCATTATATTGTTATTCTCATTTAACCTGAAAG | 2580 |
| Db | 1441 | TGCAAGCTTATGTTCCAAATATCTATCTCTAGCTCAATTGTTCCAGAGTAA | 1500 | Qy | 2581 | GAGGACTACTACATCTCCATTATGTTATTCTGTTATTGTTATTGTTATTG | 2640 |
| Qy | 1501 | AGTAGTATCAAATGCACTAACGCTCTCCTACTGGAGGTTAAAGCTTCAACATA | 1560 | Db | 2581 | GAGGACTACTACATCTCCATTATGTTATTCTGTTATTGTTATTGTTATTG | 2640 |
| Db | 1501 | AGTAGTATCAAATGCACTAACGCTCTCCTACTGGAGGTTAAAGCTTCAACATA | 1560 | Qy | 2641 | AACCTACCCAAAGTGTAGCTAGTCACTTCTGTTGCTGTTGCTGTTGCTG | 2700 |
| Qy | 1561 | CAAACGGGGAGTTTGCCTGGAATTTCTAAATGTTCTGCACTAGGTCC | 1620 | Db | 2641 | AACCTACCCAAAGTGTAGCTAGTCACTTCTGTTGCTGTTGCTGTTGCTG | 2700 |
| Db | 1561 | CAAACGGGGAGTTTGCCTGGAATTTCTAAATGTTCTGCACTAGGTCC | 1620 | Qy | 2701 | TTACCTGTTGATTAATCTGTGCTATCTGCTGTTGCTGTTGCTGTTGCTG | 2760 |
| Qy | 1621 | TCTTGTCTTAAATCTTAACTCTTAACTCTTAACTCTTAACTCTTAACTCTTAA | 1680 | Db | 2701 | TTACCTGTTGATTAATCTGTGCTATCTGCTGTTGCTGTTGCTGTTGCTG | 2760 |
| Db | 1621 | TCTTGTCTTAAATCTTAACTCTTAACTCTTAACTCTTAACTCTTAACTCTTAA | 1680 | Qy | 2761 | CTGCTCGTAAAGGGCTTAAAGGGCTTAAAGGGCTTAAAGGGCTTAAAGGG | 2820 |
| Qy | 1681 | AGAGTCAAAGGAGCTGCTGATTAATACTAGCTTAAAGCTGAGCCAGGACT | 1740 | Db | 2761 | CTGCTCGTAAAGGGCTTAAAGGGCTTAAAGGGCTTAAAGGGCTTAAAGGG | 2820 |
| Db | 1681 | AGAGTCAAAGGAGCTGCTGATTAATACTAGCTTAAAGCTGAGCCAGGACT | 1740 | Qy | 2821 | CAGAGGAAAGTCCAAACAGGACTGAACTGATAGAAAGTGTACTCTGGTGTAGTC | 2880 |
| Qy | 1741 | GTTTGTCTTAAACTCTTCACTTCTTCAACTTCTTCAACTTCTTCAACTTCTT | 1800 | Db | 2821 | CAGAGGAAAGTCCAAACAGGACTGAACTGATAGAAAGTGTACTCTGGTGTAGTC | 2880 |
| Db | 1741 | GTTTGTCTTAAACTCTTCACTTCTTCAACTTCTTCAACTTCTTCAACTTCTT | 1800 | Qy | 2881 | GCATGACATCTTGTGTTGCTGACACATACATTGGCTGCTGACACATACATT | 2940 |
| Qy | 1801 | AGTTACTAAGTCTTGTCTTCACTTCTTCAACTTCTTCAACTTCTTCAACTTCTT | 1860 | Db | 2881 | GCATGACATCTTGTGTTGCTGACACATACATTGGCTGCTGACACATACATT | 2940 |

| | | | | |
|----|------|--|------|--|
| Qy | 2941 | TATTTTGTAAGTGTGTTGATTCTGATTATATAATTCTAATATTGAAATACTCATATA | 3000 | |
| Db | 2941 | TTATTTGTAAGTGTGTTGATTCTGATTATATAATTCTAATATTGAAATACTCATATA | 3000 | |
| Qy | 3001 | TTACATGGCTTACCATTCGCCAGCAATTTCACATTGTCCTCTCATCTTGTGATG | 3060 | |
| Db | 3001 | TTACATGGCTTACCATTCGCCAGCAATTTCACATTGTCCTCTCATCTTGTGATG | 3060 | |
| Qy | 3061 | AGATGATCAGATTGGATTACTGGCTTGGAGATGATACTCATATCTATATC | 3120 | |
| Db | 3061 | AGATGATCAGATTGGATTACTGGCTTGGAGATGATACTCATATCTATATC | 3120 | |
| Qy | 3121 | TATATCTATATCTATATCTATATCTATATCTATATCTATATCTATATCTATATC | 3180 | |
| Db | 3121 | TATATCTATATCTATATCTATATCTATATCTATATCTATATCTATATC | 3180 | |
| Qy | 3181 | AAATATGTTTGTAAAGTTATAAGATTCTAGACTTTAGAACTCTGGATTGCCAAAT | 3240 | |
| Db | 3181 | AAATATGTTTGTAAAGTTATAAGATTCTAGACTTTAGAACTCTGGATTGCCAAAT | 3240 | |
| Qy | 3241 | GTAACCCCTTCCTCTACATAACCCATTTGACAAATACTTATTCATTCATC | 3300 | |
| Db | 3241 | GTAACCCCTTCCTCTACATAACCCATTTGACAAATACTTATTCATTCATC | 3300 | |
| Qy | 3301 | AAATGTTGCTGAGTCTGGCTATGAAACAGACACTGTGAAGGCCCTTGGGATATTGGC | 3360 | |
| Db | 3301 | AAATGTTGCTGAGTCTGGCTATGAAACAGACACTGTGAAGGCCCTTGGGATATTGGC | 3360 | |
| Qy | 3361 | CATGCTTGGCAAGCTTATAAGTTGCTTCATAAAAACCTTATTCTCAGTCTCATACT | 3420 | |
| Db | 3361 | CATGCTTGGCAAGCTTATAAGTTGCTTCATAAAAACCTTATTCTCAGTCTCATACT | 3420 | |
| Qy | 3421 | AACTCTGAGACTATTGCTTCAAGTACAAACTTACACTGGCTTCTATA | 3480 | |
| Db | 3421 | AACTCTGAGACTATTGCTTCAAGTACAAACTTACACTGGCTTCTATA | 3480 | |
| Qy | 3481 | TATTTGAGTAAGTAGAGCTAGAGCTAGAGCTAGAGCTAGAGCTAGAGCTAGAG | 3540 | |
| Db | 3481 | TATTTGAGTAAGTAGAGCTAGAGCTAGAGCTAGAGCTAGAGCTAGAGCTAGAG | 3540 | |
| Qy | 3541 | TGGAAATCCAACGGATAGAAAGGATGGCTCTCTGGAGAGTACGTGAGCAAAG | 3600 | |
| Db | 3541 | TGGAAATCCAACGGATAGAAAGGATGGCTCTCTGGAGAGTACGTGAGCAAAG | 3600 | |
| Qy | 3601 | GCCTGTTTGTAACTGTTCTAGTCTTGTAACTGTTCTAGTCTTGTAACTGTT | 3660 | |
| Db | 3601 | GCCTGTTTGTAACTGTTCTAGTCTTGTAACTGTTCTAGTCTTGTAACTGTT | 3660 | |
| Qy | 3661 | AATGGACAAAAACTACAAATAATGAAATGGAAACTTGGAGACAGCAAGCA | 3720 | |
| Db | 3661 | AATGGACAAAAACTACAAATAATGAAATGGAAACTTGGAGACAGCAAGCA | 3720 | |
| Qy | 3721 | TATACTATGATAAACTGTTGATTAGCTGAGAGCTTGTAACTGTTGAGCTT | 3780 | |
| Db | 3721 | TATACTATGATAAACTGTTGATTAGCTGAGAGCTTGTAACTGTTGAGCTT | 3780 | |
| Qy | 3781 | TAATGTTGCTTGTAGCTGAGCTACATCTTAACTACTCTACTAGCTTA | 3840 | |
| Db | 3781 | TAATGTTGCTTGTAGCTGAGCTACATCTTAACTACTCTACTAGCTTA | 3840 | |
| Qy | 3841 | TTTAAAAATTTTATGTTAAATGCTGAGAACTTGTAACTGTTGAGAGAG | 3900 | |
| Db | 3841 | TTTAAAAATTTTATGTTAAATGCTGAGAACTTGTAACTGTTGAGAGAG | 3900 | |
| Qy | 3901 | AGATGAAATAATTCACTTCTTACCATCTAGAAACCAATGTTGAGAGAG | 3960 | |
| Db | 3901 | AGATGAAATAATTCACTTCTTACCATCTAGAAACCAATGTTGAGAGAG | 3960 | |
| Qy | 3961 | TCCATTATTCTGCTTCTTATTCACTTACATCTTAACTGTTGAGAGAG | 4020 | |
| Db | 3961 | TCCATTATTCTGCTTCTTATTCACTTACATCTTAACTGTTGAGAGAG | 4020 | |
| Qy | 4021 | GTACATGATAACAAATGAGGCACTCTCATGTTCTCATGTTCTGCTTGTGTTAGTTA | 4080 | |
| Db | 4021 | GTACATGATAACAAATGAGGCACTCTCATGTTCTCATGTTCTGCTTGTGTTAGTTA | 4080 | |
| Qy | 4081 | ACAATATATTATGAGCATATTTCACATTTCATTTCAAAAGTTATTCTGAT | 4140 | |
| Db | 4081 | ACAATATATTATGAGCATATTTCACATTTCATTTCAAAAGTTATTCTGAT | 4140 | |
| Qy | 4141 | GGCTGTATATCACCCTACTTTGACCTTAACTGTTAGAAATACTTCTCGTGGCGTTA | 4200 | |
| Db | 4141 | GGCTGTATATCACCCTACTTTGACCTTAACTGTTAGAAATACTTCTCGTGGCGTTA | 4200 | |
| Qy | 4201 | TTTGATTTTATACTTAACTGTTAGAAATACTTCTCGTGGCTTACAGACTAGGCA | 4260 | |
| Db | 4201 | TTTGATTTTATACTTAACTGTTAGAAATACTTCTCGTGGCTTACAGACTAGGCA | 4260 | |
| Qy | 4261 | GGTGCCTGGGTCCTCAACTCCACAGATTCTGATTAACTCTGGCTGAACTAGCTAGGCA | 4320 | |
| Db | 4261 | GGTGCCTGGGTCCTCAACTCCACAGATTCTGATTAACTCTGGCTGAACTAGCTAGGCA | 4320 | |
| Qy | 4321 | TTCGAAATTCAAAAGTTCCCCAGTGATTCTAATGCTGAACTTCTGAACTGGGAACTCTTG | 4380 | |
| Db | 4321 | TTCGAAATTCAAAAGTTCCCCAGTGATTCTAATGCTGAACTTCTGAACTGGGAACTCTTG | 4380 | |
| Qy | 4381 | TAGACGGGATGATGAACTGGGTGACCTGACCTTCTGAACTTCTGAACTGGGTGACCTTCTGAACTGGGAACTCTTG | 4440 | |
| Db | 4381 | TAGACGGGATGATGAACTGGGTGACCTTCTGAACTTCTGAACTGGGAACTCTTG | 4440 | |
| Qy | 4441 | CATCTTGAGTTGCTTAAGTGTGAACTGACCATTTATAAGAACATCTGTGCAA | 4500 | |
| Db | 4441 | CATCTTGAGTTGCTTAAGTGTGAACTGACCATTTATAAGAACATCTGTGCAA | 4500 | |
| Qy | 4501 | TAATAATTATTATAGGACTCTGGTTTATGGGCTCTGCTCTAAGTTCTGAACTTGCC | 4560 | |
| Db | 4501 | TAATAATTATTATAGGACTCTGGTTTATGGGCTCTGCTCTAAGTTCTGAACTTGCC | 4560 | |
| Qy | 4561 | ATACATGCTACTAACTTAACTTAACTGACCACTTAACTGATGAACTTAACTGAA | 4620 | |
| Db | 4561 | ATACATGCTACTAACTTAACTTAACTGACCACTTAACTGATGAACTTAACTGAA | 4620 | |
| Qy | 4621 | TGTTCTGAGCTTAAACATGACCACTTAACTGATGAACTTAACTGAACTTAACTGAA | 4680 | |
| Db | 4621 | TGTTCTGAGCTTAAACATGACCACTTAACTGATGAACTTAACTGAACTTAACTGAA | 4680 | |
| Qy | 4681 | ATTATTCGCAACTTACAGGAACTGCTCATCTCTGAACTTAACTGAACTTAACTGAA | 4740 | |
| Db | 4681 | ATTATTCGCAACTTACAGGAACTGCTCATCTCTGAACTTAACTGAACTTAACTGAA | 4740 | |
| Qy | 4741 | GTATATTGCAAGATGAGAGCTGAACATGTCCTGCTGCTGCGCATA | 4800 | |
| Db | 4741 | GTATATTGCAAGATGAGAGCTGAACATGTCCTGCTGCTGCGCATA | 4800 | |
| Qy | 4801 | TGATGGCAATTAACTGTTAAAGAAATTAACTGTTAAAGAAATTAACTGCAAACT | 4860 | |
| Db | 4801 | TGATGGCAATTAACTGTTAAAGAAATTAACTGTTAAAGAAATTAACTGCAAACT | 4860 | |
| Qy | 4861 | GCCTGTTCTCTGAACTTAACTGTTAAAGAAATTAACTGTTAAAGAAATTAACTGCAACT | 4920 | |
| Db | 4861 | GCCTGTTCTCTGAACTTAACTGTTAAAGAAATTAACTGTTAAAGAAATTAACTGCAACT | 4920 | |
| Qy | 4921 | TAGGGACAAGTCTGTTGAACTTCTGCTGAGTTGAGAGAAATATTGTTGCT | 4980 | |
| Db | 4921 | TAGGGACAAGTCTGTTGAACTTCTGCTGAGTTGAGAGAAATATTGTTGCT | 4980 | |
| Qy | 5041 | ATAATAACAAATGAGCTTACCGTATGCTGTTGAACTTAACTGCAACTAA | 5100 | |
| Db | 5041 | ATAATAACAAATGAGCTTACCGTATGCTGTTGAACTTAACTGCAACTAA | 5100 | |
| Qy | 5101 | GGAAATACCAATAAGTAACAGATAACCAACAAAGGGTTACTACTAACGGCATGCGCT | 5160 | |
| Db | 5101 | GGAAATACCAATAAGTAACAGATAACCAACAAAGGGTTACTACTAACGGCATGCGCT | 5160 | |

RESULT 9
 US-10-822-873-5
 ; Sequence 5, Application US/10822873
 ; Publication No. US20040241857A1

; GENERAL INFORMATION

; APPLICANT: Henderson, D.R.
 Schuur, E.R.

; TITLE OF INVENTION: TISSUE SPECIFIC ADENOVIRAL VECTORS

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; PRIOR FILING DATE: 1995-06-27

; NUMBER OF SEQ ID NOS: 71

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; LENGTH: 5224

; TYPE: DNA

; ORGANISM: Homo sapiens

US-10-822-873-5

Query Match 100.0%; Score 5224; DB 20; Length 5224;
 Best Local Similarity 100.0%; Pred. No. 0;
 Matches 5224; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 GAATTCTAGAAATATGGGTAGGGTGCCTGCGGTAAATTCTGTTTACCCCATAGGT 60
 Db 1 GAATTCTAGAAATATGGGTAAATTCTGTTTACCCCATAGGT 60

Db 61 GAGATAGCATGGGTAAATTCTGCTTACACACATTCATAAGAATTAGG 120
 Db 61 GAGATAGCATGGGTAAATTCTGCTTACACACATTCATAAGAATTAGG 120

Qy 121 AACAGACTATGGCTGGGGACTTGAGGATTTGGACTCTCAAACTGGTGTATCT 180
 Db 121 AACAGACTATGGCTGGGGACTTGAGGATTTGGACTCTCAAACTGGTGTATCT 180

Qy 181 GTTCTATGGCTTGAACGCTGCAACTTGCACAGGGTCACTGACTTTCTCCCC 240
 Db 181 GTTCTATGGCTTGAACGCTGCAACTTGCACAGGGTCACTGACTTTCTCCCC 240

Db 241 AAGCCCAAGGPACTGCTCCTCTTCAATCTCTGTTGGGCTCTGGCTGTATCT 300
 Qy 241 AAGCCCAAGGPACTGCTCCTCTTCAATCTCTGTTGGGCTCTGGCTGTATCT 300

Db 301 GAGAAAATATAAACATTCAATAATGTTCTGGTGGATGAGATGTGTATCT 360
 Db 301 GAGAAAATATAAACATTCAATAATGTTCTGGTGGATGAGATGTGTATCT 360

Qy 361 TCATTGTGATGATGAGGACAATTAGTGTATAATCCTTAAGTACAACAACTT 420
 Db 361 TCATTGTGATGATGAGGACAATTAGTGTATAATCCTTAAGTACAACAACTT 420

Qy 421 GAGGGTAGGGTGGTACTATCAATTCTTATTATAAGGATACTTCTTATTATA 480
 Qy 421 GAGGGTAGGGTGGTACTATCAATTCTTATTATAAGGATACTTCTTATTATA 480

Db 1321 ACAACTCTGGTCAAGCTCCTCTTATGCTGCTCTGGAAATTGCTGTTCTTCAT 1380
 Db 1321 ACAACTCTGGTCAAGCTCCTCTTATGCTGCTCTGGAAATTGCTGTTCTTCAT 1380

Qy 1381 GTTCTCTTCTGCTCTGGCTCATCTGCTATCTTCTGCTCAAACTCACATGTC 1440
 Db 1381 GTTCTCTTCTGCTCTGGCTCATCTGCTATCTTCTGCTCAAACTCACATGTC 1440

Qy 1441 TGCAGCTPATGATGCCAAATACTTCTGCTCAATCTGCTCAAACTCACATGTC 1500
 Db 1441 TGCAGCTPATGATGCCAAATACTTCTGCTCAATCTGCTCAAACTCACATGTC 1500

Qy 1501 AGTAACTTCAACATCAACCTCTCCACTTGGGCTTAAGAACCTTCCACACATA 1560
 Db 1501 AGTAACTTCAACATCAACCTCTCCACTTGGGCTTAAGAACCTTCCACACATA 1560

| | | | |
|----|------|---|------|
| Qy | 1551 | CAACCGGGAGTTTCCTAAATACTGCTGGATGTTGCCTGAGCATGGGTCC | 1620 |
| Db | 1551 | CAACCGGGAGTTGCCTGAGCATGGGTCC | 1620 |
| Qy | 1621 | TCTGTTCCTTAAATCTAAATTACTTTAGGCCAGTGCTCCTAACCTATGGGAGATG | 1680 |
| Db | 1621 | TCTGTTCCTTAAATCTAAATTACTTTAGGCCAGTGCTCCTAACCTATGGGAGATG | 1680 |
| Qy | 1681 | AGGTGAAAGGGAGCTGTAAACTTAATCACTAAGTCACTGCTAACCTATGGGAGCT | 1740 |
| Db | 1681 | AGGTGAAAGGGAGCTGTAAACTTAATCACTAAGTCACTGCTAACCTATGGGAGCT | 1740 |
| Qy | 1741 | GTTGGTAAACTGTCACTTAACTAACTAACTAACTAACTAACTAACTAACTAACT | 1800 |
| Db | 1741 | GTTGGTAAACTGTCACTTAACTAACTAACTAACTAACTAACTAACTAACTAACT | 1800 |
| Qy | 1801 | AGCTTAAAGTCCTTGAACCTTATCTCATTCTACATCAGCTTATCCAGGCCACTTAT | 1860 |
| Db | 1801 | AGTAACTAAGTCCTTGAACCTTATCTCATTCTACATCAGCTTATCCAGGCCACTTAT | 1860 |
| Qy | 1861 | TTCACAGTATTGCGAAAACCTCCTAACCTGCTCCTTATCATAGCTPATCCCCTT | 1920 |
| Db | 1861 | TTCACAGTATTGCGAAAACCTCCTAACCTGCTCCTTATCATAGCTPATCCCCTT | 1920 |
| Qy | 1921 | TGAAACAAAGGAGCACGTTCAAATAAAATGAAATGATTGTTATTAGTCTCCCTTTGGT | 1980 |
| Db | 1921 | TGAAACAAAGGAGCACGTTCAAATAAAATGAAATGATTGTTATTAGTCTCCCTTTGGT | 1980 |
| Qy | 1981 | CTATAATAGTCCTCAGGGAGTTATAACTCCATTAAAGCTTGTAGATGTCGCCCT | 2040 |
| Db | 1981 | CTATAATAGTCCTCAGGGAGTTATAACTCCATTAAAGCTTGTAGATGTCGCCCT | 2040 |
| Qy | 2041 | TGCCAACACTTGCAGGAAATTCCCAATATCTAGTATTTCCTACTATAACTTGTGCCCT | 2100 |
| Db | 2041 | TGCCAACACTTGCAGGAAATTCCCAATATCTAGTATTTCCTACTATAACTTGTGCCCT | 2100 |
| Qy | 2101 | TTCAAACTGCAATTTCCTCATCCCTAACCTGCTAACCTGCTAACCTGCTAACCTGCT | 2160 |
| Db | 2101 | TTCAAACTGCAATTTCCTCATCCCTAACCTGCTAACCTGCTAACCTGCTAACCTGCT | 2160 |
| Qy | 2161 | TTCACCCACCTTTACATTTCCTGGAAACACTATACCTCCCTCTICATTGGCCACC | 2220 |
| Db | 2161 | TTCACCCACCTTTACATTTCCTGGAAACACTATACCTCCCTCTICATTGGCCACC | 2220 |
| Qy | 2221 | TCTTAATTCTCTTCAGATCTCCATGAGATGTTACTTCCTCAGGAAGGCCCTATCTGACC | 2280 |
| Db | 2221 | TCTTAATTCTCTTCAGATCTCCATGAGATGTTACTTCCTCAGGAAGGCCCTATCTGACC | 2280 |
| Qy | 2281 | CCTCCAAGATGTCATGAGTTCTCTTTCACTCTAACTAACTACAGATCCATCACCA | 2340 |
| Db | 2281 | CCTCCAAGATGTCATGAGTTCTCTTTCACTCTAACTACAGATCCATCACCA | 2340 |
| Qy | 2341 | TGTTGTGATTACTGATGACTATTCTGTTCTGATTGGGAGTAACTTCTCATTAACTGAGC | 2400 |
| Db | 2341 | TGTTGTGATTACTGATGACTATTCTGTTCTGATTGGGAGTAACTTCTCATTAACTGAGC | 2400 |
| Qy | 2401 | TACATGTCGCCCTCTCTGTCGTTGCTGATTAACTCCATCCAAACAGTCCTGGATGCC | 2460 |
| Db | 2401 | TACATGTCGCCCTCTCTGTCGTTGCTGATTAACTCCATCCAAACAGTCCTGGATGCC | 2460 |
| Qy | 2461 | GACTTAACTTTATGATGATGATAATAATAAACCCATCTATGGAGTCACTTGTGCA | 2520 |
| Db | 2461 | GACTTAACTTTATGATGATGATAATAATAAACCCATCTATGGAGTCACTTGTGCA | 2520 |
| Qy | 2521 | AGACCCGGTTCTGAGGCAATTATGATTAACTTCTCATTAACTGAGTC | 2580 |
| Db | 2521 | AGACCCGGTTCTGAGGCAATTATGATTAACTTCTCATTAACTGAGTC | 2580 |
| Qy | 2581 | GAGTCACTATCACTATCCCTTATTTATGTTGATAAGATAAGCCAGAAATGAAATT | 2640 |
| Db | 2581 | GAGTCACTATCACTATCCCTTATTTATGTTGATAAGATAAGCCAGAAATGAAATT | 2640 |
| Qy | 2641 | AACTCACCCAAAGTCATGTCAGGCAAAAACTAAACGTTCCCCACT | 2700 |

| | | | | | | | |
|----|------|--|------|-----------|------|--|------|
| Db | 3721 | TATACTATGATAAATGTTGTATTGCTGGAAAGTGCTACTGACGGTAACCCCTTTGTT | 3780 | Qy | 4861 | GCTGTCCTTCTTATGGCTCATTAACCTTGTGCACT | 4920 |
| Qy | 3781 | TAATATGTCGCCCTTAGTPGCTGCACTGATGCTTAACTGTTAAGTACTGTACTTACGTTA | 3840 | Db | 4861 | GCTGTCCTTCTTATGGCTCATTAACCTTGTGCACT | 4920 |
| Db | 3781 | TAATATGTCGCCCTAGTPGCTGCACTGATGCTTAACTGTTAAGTACTGTACTTACGTTA | 3840 | Qy | 4921 | TAGGACAAGTCATGCTTGAATATTCTGTAGTTGAGGAAATATTGTGTTATTGAGAATTTG | 4980 |
| Qy | 3841 | TTAAAAAAATTGTTAATTGCTTAACTGCTTCACTGAGTTTCATGAAAGAATTTGAGAG | 3900 | Db | 4921 | TAGGACAAGTCATGCTTGAATATTCTGTAGTTGAGGAAATTTGAGAATTTG | 4980 |
| Db | 3841 | TTAAAAAAATTGTTAATTGCTTAACTGCTTCACTGAGTTTCATGAAAGAATTTGAGAG | 3900 | Qy | 4981 | AAATAAAATTAAGTTGCAACTTTCTGCACTTTCTGCTGTCCTGTCCTGAAAC | 5040 |
| Qy | 3901 | AGATAGATTAAATTCACTTACCTTACATTPGAGAACCCAATGTTAAACTGTG | 3960 | Db | 4981 | AAATAAAATTAAGTTGCAACTTTCTGCACTTTCTGCTGTCCTGTCCTGAAAC | 5040 |
| Db | 3901 | AGATAGATTAAATTCACTTACATTPGAGAACCCAATGTTAAACTGTG | 3960 | Qy | 5041 | ATAAALATACAAAACCGPATGCTTTAAATTGTCACATTTCAACCTAA | 5100 |
| Qy | 3961 | TCCATTATTCTGCTTATTCAACATTTTTACAGGGGGAAATACAGGGAG | 4020 | Db | 5041 | ATAAALATACAAAACCGPATGCTTTAAATTGTCACATTTCAACCTAA | 5100 |
| Db | 3961 | TCCATTATTCTGCTTATTCAACATTTTTACAGGGGGAAATACAGGGAG | 4020 | Qy | 5101 | GGAAATACCATTAAGTAACAGATAACCAAAAGGTTACTAGTTAACAGGCA | 5160 |
| Qy | 4021 | GTACAATGATACACAAATGAGAACACTCCATGATTGTTCTGTTTCAGTTA | 4080 | Db | 5101 | GGAAATACCATTAAGTAACAGATAACCAAAAGGTTACTAGTTAACAGGCA | 5160 |
| Db | 4021 | GTACAATGATACACAAATGAGAACACTCCATGATTGTTCTGTTTCAGTTA | 4080 | Qy | 5161 | GAAAGAGTATAAAAGAATTTCAGCATGTTCCATATGTCCTTCACCAACTG | 5220 |
| Qy | 4081 | ACATATATTAGCATATTTCATTCAATTATCATTGAAAGTTATTGAT | 4140 | Db | 5161 | GAAAGAGTATAAAAGAATTTCAGCATGTTCCATATGTCCTTCACCAACTG | 5220 |
| Db | 4081 | ACATATATTAGCATATTTCATTCAATTATCATTGAAAGTTATTGAT | 4140 | Qy | 5221 | AACA | 5224 |
| Qy | 4141 | GCCTGATATCACCTPACTTTATGATGACCAATTAAATTCTGGTTGGTTA | 4200 | Db | 5221 | AACA | 5224 |
| Db | 4141 | GCCTGATATCACCTPACTTTATGATGACCAATTAAATTCTGGTTGGTTA | 4200 | RESULT 10 | | | |
| Qy | 4201 | TTTGATTTTATATCTTACCTTACGTTAGATAAAGCCTGAAATACT | 4260 | | | | |
| Db | 4201 | TTTGATTTTATATCTTACCTTACGTTAGATAAAGCCTGAAATACT | 4260 | | | | |
| Qy | 4261 | GGTGCCTGGCTCACTCCAGATTCTGTTAACTGCTGCTGGTTACGACTAGGA | 4320 | | | | |
| Db | 4261 | GGTGCCTGGCTCACTCCAGATTCTGTTAACTGCTGCTGGTTACGACTAGGA | 4320 | | | | |
| Qy | 4321 | TTGGGAAATCAAAAAGTTCCCTCACTGTTAACTGCTGCTGGTTACGACTAGGA | 4380 | | | | |
| Db | 4321 | TTGGGAAATCAAAAAGTTCCCTCACTGTTAACTGCTGCTGGTTACGACTAGGA | 4380 | | | | |
| Qy | 4381 | TAGACAGGATGATAGGAGTGAACCTTACATCATTTAGTATTACATCAT | 4440 | | | | |
| Db | 4381 | TAGACAGGATGATAGGAGTGAACCTTACATCATTTAGTATTACATCAT | 4440 | | | | |
| Qy | 4441 | CATCTGAGTCTAAGTGTGATGCTTAACTGAGCATGCAA | 4500 | | | | |
| Db | 4441 | CATCTGAGTCTAAGTGTGATGCTTAACTGAGCATGCAA | 4500 | | | | |
| Qy | 4501 | TAAAATTATTAGGACTTAACTTAAATGTTACCTTATTGACATACATTAAGTCAAAG | 4620 | | | | |
| Db | 4501 | TAAAATTATTAGGACTTAACTTAAATGTTACCTTATTGACATACATTAAGTCAAAG | 4620 | | | | |
| Qy | 4561 | ATACATGCAACTTAACTTAAATGTTACCTTATTGACATACATTAAGTCAAAG | 4620 | | | | |
| Db | 4561 | ATACATGCAACTTAACTTAAATGTTACCTTATTGACATACATTAAGTCAAAG | 4620 | | | | |
| Qy | 4621 | TGTTTCAGCTGAGCTAACATGCACTGACCTGACCTTAACTGAAATTT | 4680 | | | | |
| Db | 4621 | TGTTTCAGCTGAGCTAACATGCACTGACCTGACCTTAACTGAAATTT | 4680 | | | | |
| Qy | 4681 | ATTATCTGCAACTTAACTGACCTGACCTTAACTGAAATTTGACTTGAGACTATT | 4740 | | | | |
| Db | 4681 | ATTATCTGCAACTTAACTGACCTGACCTTAACTGAAATTTGACTTGAGACTATT | 4740 | | | | |
| Qy | 4741 | GTTATATTGCAAGTGAAGTGAAGTCTGCTGCTGCTGCTGCTGCTGCTG | 4800 | | | | |
| Db | 4741 | GTTATATTGCAAGTGAAGTGAAGTCTGCTGCTGCTGCTGCTGCTGCTG | 4800 | | | | |
| Qy | 4801 | TGATAGGCAATTAAATGTTAAAGAATTAAATGTTAGTGAATTGCTACCAACT | 4860 | | | | |
| Db | 4801 | TGATAGGCAATTAAATGTTAAAGAATTAAATGTTAGTGAATTGCTACCAACT | 4860 | | | | |

| id | seq | label |
|------|---|-------|
| 121 | AACAGACTATEGGCTGGAGGACTTTAGGATGTCAGGTCTCATAACACTTGGGTGTATCT | 180 |
| 181 | GTTCTATGGGTTGGTTAGCTGGCAACTTGGAAAGGGTCACTGACTTCTCC | 240 |
| 181 | GTTCTATGGGTTGGTTAGCTGGCAACTTGGAAAGGGTCACTGACTTCTCC | 240 |
| 241 | AAGCCCAGGTACTGCTCTTTACATCTGTTGGGGCTCTGGGCTGAATACT | 300 |
| 241 | AAGCCCAGGTACTGCTCTTTACATCTGTTGGGCCTGAATACTGACTTCTCC | 300 |
| 301 | GAGAAATAAACAATTCAAACTTCTGTTGGAGATGAGATGAGATGCTCAT | 360 |
| 301 | GAGAAATAAACAATTCAAACTTCTGTTGGAGATGAGATGAGATGCTCAT | 360 |
| 361 | TCACTGTATCAATGATGATGATGAACTTGGTATAAATCCTTAGACAACT | 420 |
| 361 | TCACTGTATCAATGATGATGATGAACTTGGTATAAATCCTTAGACAACT | 420 |
| 421 | GAGGTAGGGGGTACTATCAATTCTATTAAAGATACTTATTTATTAA | 480 |
| 421 | GAGGTAGGGGGTACTATCAATTCTATTAAAGATACTTATTTATTAA | 480 |
| 481 | TGCTCTGACAAATGTTGCGGACCAAGGATCAGAAAGTGAGTCCTTGAAATT | 540 |
| 481 | TGCTTGACAAATGTTGCGGACCAAGGATCAGAAAGTGAGTCCTTGAAATT | 540 |
| 541 | AAGAAGTTAATGGTCCAGGAAATAATCATAGCTTACAANTGACTGATACTCAA | 600 |
| 541 | AAGAAGTTAATGGTCCAGGAAATAATCATAGCTTACAANTGACTGATACTCAA | 600 |
| 601 | ACAAGAGGTTCCATGAGAAATAATCTGAAAGGTTAAATAGGTCAGGGTGAGGG | 660 |
| 601 | ACAAGAGGTTCCATGAGAAATAATCTGAAAGGTTAAATAGGTCAGGGTGAGGG | 660 |
| 661 | CTCTCTCTGCTGAGACTAATCGAAATAACATTCAGGGATAATTCTGAACT | 720 |
| 661 | CTCTCTCTGCTGAGACTAATCGAAATAACATTCAGGGATAATTCTGAACT | 720 |
| 721 | TAAGGGTTGGTACATTGGTCAAGGATGTTGAGAAAGGAGGTGAAATTGAAAC | 780 |
| 721 | TAAGGGTTGGTACATTGGTCAAGGATGTTGAGAAAGGAGGTGAAATTGAAAC | 780 |
| 781 | ATTTCACTAACCACCAATTCCAACAAACAAAAATTGAAAGAACTCTGAAACA | 840 |
| 781 | ATTTCACTAACCACCAATTCCAACAAACAAAAATTGAAAGAACTCTGAAACA | 840 |
| 841 | GTGAGATAAGGAGAGGAATTTCACACCCACGTATGCCAACNGCTGAGAA | 900 |
| 841 | GTGAGATAAGGAGAGGAATTTCACACCCACGTATGCCAACNGCTGAGAA | 900 |
| 901 | GTATATATCTATAATTAACTAACTATCATGCTTATAATGATAATAATTACTGCAATT | 960 |
| 901 | GTATATATCTATAATTAACTAACTATCATGCTTATAATGATAATAATTACTGCAATT | 960 |
| 1021 | ACTTGAGGGATAGTCAGTCATTTCATGATAATGAGAAATTAAACAGATCAATT | 1080 |
| 1021 | ACTTGAGGGATAGTCAGTCATTTCATGATAATGAGAAATTAAACAGATCAATT | 1080 |
| 1081 | TTGCTCTGTCATACAGCTTAATTGACCTAAAGCAATTAGGTTAAATTGTTGAAAT | 1140 |
| 1081 | TTGCTCTGTCATACAGCTTAATTGACCTAAAGCAATTAGGTTAAATTGTTGAAAT | 1140 |
| 1141 | CTTCTTAATCCAACTTCAGTTACTGTTCACTGTTCTGCTGCTGACTCTGT | 1260 |
| 1141 | CTTCTTAATCCAACTTCAGTTACTGTTCACTGTTCTGCTGACTCTGT | 1260 |
| 1201 | TATGAAAAGTAAACGAAATTACATCANTCAGAAAGCATGCTGACTCTGT | 1260 |
| 1201 | TATGAAAAGTAAACGAAATTACATCANTCAGAAAGCATGCTGACTCTGT | 1260 |

Db 4501 TAAAATTATTAAGGGACTGGTTTATTAGGGCTTGTGCTCTAAGTTCTATGTTAAGCC 4560
 Qy 4561 ATACATCGCATACTAAATCACTTTAAATGACCTTATTGACATACATAATTAGTGAAGG 4620
 Db 4561 ATACATCGCATACTAAATCACTTTAAATGACCTTATTGACATACATAATTAGTGAAGG 4620
 Qy 4621 TGTTCTGAGCTAACATGACAGCATAATTACGAAATTGATAATTGAAATTGTT 4680
 Db 4621 TGTTCTGAGCTAACATGACAGCATAATTACGAAATTGATAATTGAAATTGTT 4680
 Qy 4681 ATTATTCGCAACTTGGGACAAGTCATCTCAGAAATTGTTGACTTGTGAGGTATT 4740
 Db 4681 ATTATTCGCAACTTGGGACAAGTCATCTCAGAAATTGTTGACTTGTGAGGTATT 4740
 Qy 4741 GTTATATTGAGATGAGGTGAAAGTCGAACTTGTCAGACATGCTGTTCCGCAAA 4800
 Db 4741 GTTATATTGAGATGAGGTGAAAGTCGAACTTGTCAGACATGCTGTTCCGCAAA 4800
 Qy 4801 TGATAGGCAATTAAAGATAATTGATTGATGATTGATGAACTTCAACAAATCT 4860
 Db 4801 TGATAGGCAATTAAAGATAATTGATTGATGATTGATGAACTTCAACAAATCT 4860
 Qy 4861 GCTGCTTCTTCTTATGGCCTCATTAACCTTAATTGAGAAATTAAATTCTGCAACT 4920
 Db 4861 GCTGCTTCTTCTTATGGCCTCATTAACCTTAATTGAGAAATTAAATTCTGCAACT 4920
 Qy 4921 TAGGCCAGTCATCTGAAATTGATTCTAGTTGAGGAGATTGTTGATATTGCT 4980
 Db 4921 TAGGCCAGTCATCTGAAATTGATTCTAGTTGAGGAGATTGTTGATATTGCT 4980
 Qy 4981 AAAATAAAATAAGTTGCAAGTCTTCTGCCCCAAATGTCCTGCTGCTGCAC 5040
 Db 4981 AAAATAAAATAAGTTGCAAGTCTTCTGCCCCAAATGTCCTGCTGCTGCAC 5040
 Qy 5041 ATAALATAACATAACCGTATGCTGTTGAAATTCTGCTTGTGAGGATTCACCT 5100
 Db 5041 ATAALATAACATAACCGTATGCTGTTGAAATTCTGCTTGTGAGGATTCACCT 5100
 Qy 5101 GGAAATACATATACTACAGATAACCCAAATGCTGTTGAAATTCTGCTTGTGCT 5160
 Db 5101 GGAAATACATATACTACAGATAACCCAAATGCTGTTGAAATTCTGCTTGTGCT 5160
 Qy 5161 GAAAGAGTTAAAGAAATTCAAGATGATTTCAGATGATTTCAGATGATTTCAG 5220
 Db 5161 GAAAGAGTTAAAGAAATTCAAGATGATTTCAGATGATTTCAGATGATTTCAG 5220
 Qy 5221 AACAA 5224
 Db 5221 AACAA 5224
 Qy 5221 AACAA 5224
 Db 5221 AACAA 5224

RESULT 11
 US-10-691-045-16
 Sequence 16, Application US/10691045
 Publication No. US20010146489A1
 GENERAL INFORMATION:
 APPLICANT: Yu, De-Chao
 APPLICANT: Li, Yuanhao
 APPLICANT: Henderson, Daniel R.
 TITLE OF INVENTION: CELL-SPECIFIC ADENOVIRUS VECTORS
 TITLE OF INVENTION: COMPRISING AN INTERNAL RIBOSOME ENTRY SITE
 CURRENT APPLICATION NUMBER: US/10/691,045
 CURRENT FILING DATE: 2003-10-21
 PRIOR APPLICATION NUMBER: US/09/814,351
 PRIOR FILING DATE: 2001-03-21
 PRIOR APPLICATION NUMBER: 60/192,156
 PRIOR FILING DATE: 2000-03-24
 NUMBER OF SEQ ID NOS: 35
 SOFTWARE: FastSEQ for Windows Version 4.0
 SEQ ID NO: 16
 LENGTH: 5224
 TYPE: DNA

; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: AlphaFP-TRE
; US-10-691-045-16
Query Match 100.0%; Score 5224; DB 22;
Best Local Similarity 100.0%; Pred. No. 0;
Matches 5224; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
1 GAATCTCTAGAAATAAGGGTAACTGGGTGTTGGGTAAATTCTCTTTCACCCCTAGGT 60
1 GAATCTCTAGAAATAAGGGTAACTGGGTGTTGGGTAAATTCTCTTTCACCCCTAGGT 60
61 GAGTAGGATTGGTTAAATGGCTTCAACACACATCACATTCAAGATAATTAGG 120
61 GAGTAGGATTGGCTTAAATGGCTTCAACACACATCACATTCAAGATAATTAGG 120
121 AACAGACATGGCTGAGGACTTGGGGCTGCTCTCATACACTPTGGGTATCT 180
121 AACAGACATGGCTGAGGACTTGGGGCTGCTCTCATACACTPTGGGTATCT 180
181 GTTCATGGGCTTAAAGCTGGAACTTGGGACTTGGGGCTTGGAAATATCT 240
181 GTTCATGGGCTTAAAGCTGGAACTTGGGACTTGGGCTTGGAAATATCT 240
241 AAGCCAAAGTGTCTGCTCTTCAATCTGTTGGGCTCTGGGCTTGGATATCT 300
241 AAGCCAAAGTGTCTGCTCTTCAATCTGTTGGGCTCTGGGCTTGGATATCT 300
301 GAGAAATAAAACATTCAATATGCTGCTGCTGAGGATGAGGATGTCAT 360
301 GAGAAATAAAACATTCAATATGCTGCTGCTGAGGATGAGGATGTCAT 360
361 TCATTGTATCAATGAAATTGAGGACATTAGTGATAATCTTACAACT 420
361 TCATTGTATCAATGAAATTGAGGACATTAGTGATAATCTTACAACT 420
421 GAGGGTAGGGCTGCTATTCAATTCTPATTTAAAGATACTTCTATTATA 480
421 GAGGGTAGGGCTGCTATTCAATTCTPATTTAAAGATACTTCTATTATA 480
481 TGCTTGACAAATGTTGTTGGGACCAAGGAAATCAAAGATGACTCTTGTGAACT 540
481 TGCTTGACAAATGTTGTTGGGACCAAGGAAATCAAAGATGACTCTTGTGAACT 540
541 AAGAGTTATGGTCAGGAATTATACTAGCTTACAAATTGACTATCCATCA 600
541 AAGAGTTATGGTCAGGAATTATACTAGCTTACAAATTGACTATCCATCA 600
601 ACAGAGGTTCCATGAGAAATTCTGAAAGGTTAAATGTTGCAAAAGTGAGGG 660
601 ACAGAGGTTCCATGAGAAATTCTGAAAGGTTAAATGTTGCAAAAGTGAGGG 660
661 CTCTCTCTGAGGACTAACTGAAATACTTCACTGGGATATATTGTAATGACCT 720
661 CTCTCTCTGAGGACTAACTGAAATACTTCACTGGGATATATTGTAATGACCT 720
721 TAACGGTTGGGTACATTGTTGACATTGATGGAGGAGGAGTGAATTGGAAAC 780
721 TAACGGTTGGGTACATTGTTGACATTGATGGAGGAGGAGTGAATTGGAAAC 780
781 ATTTCACAACTAAACCAACCCAAATCCAACAAACAAATAAGATAAAAGAACTCAGAAACA 840
781 ATTTCACAACTAAACCAACCCAAATCCAACAAACAAATAAGATAAAAGAACTCAGAAACA 840
841 GTGAGATAAGGAGGAATTCTCACACCCACGTTAGCTCAACTGCTCTGAAAGA 900
841 GTGAGATAAGGAGGAATTCTCACACCCACGTTAGCTCAACTGCTCTGAAAGA 900
901 GTATATCTAAATTAACTTAAACTAACATGCTTAAATGATAATACTGCTCATT 960
901 GTATATCTAAATTAACTTAAACTAACATGCTTAAATGATAATACTGCTCATT 960

| | | | |
|----|------|---|------|
| Db | 2701 | GATCGATATTGATGAACTGGTGTGATAATATATTTGTTGATGTTGATGTTGATGATA | 2760 |
| Qy | 2941 | TATTTGTAGTGTGTGTGTTATTATTTATATATTTGAAATAGTCATAAT | 3000 |
| Db | 2761 | TATTTGTAGTGTGTGTGTTATTATTTATATATTTGAAATAGTCATAAT | 2820 |
| Qy | 3001 | TTACTAAGCCCTACCAATTGCGAGGATTTCATTTGCCCCCTTAATCTTGTG | 3060 |
| Db | 2821 | TTATTAAAGGTTTATTGGTAGTTTATTGGTAGTTTATTGGTAGTTTGTG | 2880 |
| Qy | 3061 | AGATGATCAGATTGGATTACTGGCCTTGAAAGTAGATACTACATCTATCATC | 3120 |
| Db | 2881 | AGATGATCAGATTGGATTACTGGCCTTGAAAGTAGATACTACATCTATCATC | 2928 |
| Qy | 3121 | TATATCTATACCTATATCTATATCTATATCTATATCTATATCTATATCTAT | 3180 |
| Db | 2929 | TATATCTATATTTATATTTATATTTATATTTATATTTATATTTATATCTAT | 2988 |
| Qy | 3181 | AAATAGCTTTGTAAGCTTAAGATTCAAGCTTTAGATCTGGATTGCGCAAT | 3240 |
| Db | 2989 | AAATAGCTTTGTAAGCTTAAGATTCAAGCTTTAGATCTGGATTGCGCAAT | 3048 |
| Qy | 3241 | GTAACCCCTTCTCATTAACCATGTTGAAACAATACATTCTATCATC | 3300 |
| Db | 3049 | GTAATTTTTTTTATTTATTTATTTATTTATTTATTTATTTATTTATTTAT | 3108 |
| Qy | 3301 | AAATGTTGCTGAGTCTGGCTATGAAACCCTTGGGATATTGCC | 3360 |
| Db | 3109 | AAATGTTGCTGAGTCTGGCTATGAAACCCTTGGGATATTGCC | 3168 |
| Qy | 3361 | CATGCTTGGCAGCCTATACTGTTGCTTATAACTCTATTGCTCTCATACT | 3420 |
| Db | 3169 | TATGTTGGCTAAGTTTATACTGTTGCTTATAACTCTATTGCTCTCAT | 3228 |
| Qy | 3421 | AATACTCTACGACTATTGCTTTCTGTTCTATGAAACTCTGCTTCATA | 3480 |
| Db | 3229 | ATATTTGATTTGTTTATTTGTTTATTTGTTTATTTGTTTATTTGTT | 3288 |
| Qy | 3481 | TATTGAGTAAGTGCCTTGAGGAAGACTAGAAGTAACTATCATPATCC | 3540 |
| Db | 3289 | TATTGAGTAAGTGCCTTGAGGAAGACTAGAAGTAACTATCATPATCC | 3348 |
| Qy | 3541 | TGGAATCCTAACGGGATAGACAGGATGGGTGACTCTTCTGGAGTAGCAAG | 3600 |
| Db | 3349 | TGGAATTTAACGGGATAGACAGGATGGGTGACTCTTCTGGAGTAGCAAG | 3408 |
| Qy | 3601 | GCCTGTTTGTAACTGTTCTTACATGTTCTTACATGAACTTGGAGAGACCAAA | 3660 |
| Db | 3409 | GTGTTGTTGTTATGTTTACATGAACTTGGAGAGACCAAA | 3468 |
| Qy | 3661 | ATGGCAAAACAAACTAAACATGAATGGGAATTTGACTCTGTTGAGCTTGT | 3720 |
| Db | 3469 | ATGGCAAAACAAATTAAATGAATGGGAATTTGACTCTGTTGAGCTTGT | 3528 |
| Qy | 3781 | TAACTGTGTCCTCTAGCTAGCTGCACTATGATCTTAACTGTTGAGCTT | 3840 |
| Db | 3589 | TAATGTGTCCTCTAGCTAGCTGCACTATGATCTTAACTGTTGAGCTT | 3648 |
| Qy | 3841 | TTAAAAAAATTATGTTAAATTGCACTAGCTGCTCTTCAATGAAAGTTGAGAG | 3900 |
| Db | 3529 | TTAAAAAAATTATGTTAAATTGCACTAGCTGCTCTTCAATGAAAGTTGAGAG | 3708 |
| Qy | 3901 | AGATGAAATTAAATTGCACTAGCTGCTCTTCAATGAAAGTTGAGAG | 3960 |
| Db | 3709 | AGATGAAATTAAATTGCACTAGCTGCTCTTCAATGAAAGTTGAGAG | 3768 |
| Qy | 3961 | TCCATTATTCTGCTCTTCACTGCTTAACTGCTTAACTGCTTAACTGCTTAA | 4020 |
| Db | 3769 | TTATTATTGTTTATTAACTGCTTAACTGCTTAACTGCTTAACTGCTTAA | 3828 |

Sequence 59 , Application US/10172086
 Publication No. US20030113750A1
 GENERAL INFORMATION:
 APPLICANT: Epigenomics AG
 TITLE OF INVENTION: Method and nucleic acids for the differentiation
 of prostate tumors
 FILE REFERENCE:
 CURRENT APPLICATION NUMBER: US/10/172,086
 CURRENT FILING DATE: 2002-06-13
 NUMBER OF SBQ ID NOS: 116
 SEQ ID NO 59
 TYPE: DNA
 ORGANISM: Artificial Sequence
 FEATURE:
 OTHER INFORMATION: chemically treated genomic DNA (Homo sapiens)
 US-10-172-086-59

| Query | Match | Score | Length |
|-------|-----------------------|--|---------------------|
| Db | Best Local Similarity | 84.3% | 7001; |
| Db | Matches | 986 | Pred. No. 2.6e-174; |
| Db | Conservative | 0 | Mismatches 184; |
| Db | Indels | 0 | Gaps 0; |
| Qy | 4055 | TATTGTTTGGCTCTGTTTCAGTTACAATATGAGCATATTCCATTTCATTAA | 4114 |
| Db | 1 | TATTGTTTGGTTTAGTTATAATATGAGTATTTTATTGATGTATTAA | 60 |
| Qy | 4115 | ATATTCCTCCACAAAGTTATTTGANGGCSTATAATACCCCTACTTTGATGTACCAT | 4174 |
| Db | 61 | ATATTTTTATAAAGTTATTGTGTTGTTGTTATTTATGAGTATTTGATGTATTAA | 120 |
| Qy | 4175 | ATTAAATTATTTCCTCTGTGGGTATTGTATTGTTATAATCTCCCTTAGATAATGAA | 4234 |
| Db | 121 | ATTAAATTATTTCCTGTGGGTATTGTATTGTTATAATCTCCCTTAGATAATGAA | 180 |
| Qy | 4235 | ACACCTTGAGCTTACAGATACTGTTAGAAATACTGTGCTCGGTCTCACTCCACAGTTCTGATT | 4294 |
| Db | 181 | ATATTTGTGAGTTTGTGAAATTTGTTGGTTTAATTTTATGATTTGATTTGATTT | 240 |
| Qy | 4295 | AACCTGGCTCTGGTTACAGACTAGCCATTAGGTTAGGAACTTCAAAGGTTCCCAGCTGATCTAA | 4354 |
| Db | 241 | AATTGGTTGGTTAGTTAGATTAGTTAGGTTAGGAAATTAAAAGTTTTACTGATTTAA | 300 |
| Qy | 4355 | TGTGTGGCCAGATCGGAAACCTCTAGACGGGTGATAGGGGTGAGCCACTCTAG | 4414 |
| Db | 301 | TGTGTGGCCAGATCGGAAACCTCTAGACGGGTGATAGGGGTGAGCCACTCTAG | 360 |
| Qy | 4415 | CATCCATCATTAGPATAACATCATCATGCAAAATAATTATAGGACTCTGGTTATTGGGCTT | 4534 |
| Db | 421 | TTATTTTATAAGATAATATGTTGTAATAAATTATAGGATTGGTTATTGGGCTT | 480 |
| Qy | 4535 | GTGCTCTAAAGACACATGTGCAAAATAAAATTATAGGACTCTGGTTATTGGGCTT | 4594 |
| Db | 361 | TATTGTTTAAAGTTGTTAGTTGTTAGTTGTTAGTTGATGTTAGTTGATGTTGAT | 420 |
| Qy | 4475 | CCACCTTATAAAAGACACATGTGCAAAATAAAATTATAGGACTCTGGTTATTGGGCTT | 4534 |
| Db | 421 | TTATTTTATAAGATAATATGTTGTAATAAATTATAGGATTGGTTATTGGGCTT | 480 |
| Qy | 4595 | TATTGACATACATTTAGTGAAGAGTGTGTTCTGGCTAACAAATCACGCTAAATTAC | 4654 |
| Db | 481 | GTGCTCTAAAGTTGTTAGTTGTTAGTTGTTAGTTGATGTTGTTGATGTTGAT | 540 |
| Qy | 4655 | AAGCATGATAATTGAAATTATTGTTGAGGTATATTGCAAGTGAAGCTCATCTCT | 4714 |
| Db | 601 | AAGTATGATAATTGAAATTGTTGAGGTATATTGTAAGTGAAGTTGAAATTGTT | 660 |
| Qy | 4715 | GAATTGTTGACTTGTGAGGTATATTGCAAGTGAAGCTGAAATGGCT | 4774 |
| Db | 661 | GAATTGTTGATTTGTTGAGGTATATTGTAAGTGAAGTTGAAATTGTT | 720 |
| Qy | 4775 | AGACATGTCCTGTCGCTGGCTGGCATATGATAGGGCTTAATAGTTAAAGAATTAAATGT | 4834 |
| Db | 721 | AGATAATGTTGTTGTTGTTGTTGATAGTTAATGATGTTAATGATGTTAATGAT | 780 |

RESULT 15 -
 US-10-172-086-59

| | | | |
|----|------|--|------|
| Qy | 4835 | ATTTAGATGAAATTGCCATAACCAATCTGCCTCTTCTTATGGCTTCATTAACCTTAAT ^{NT} | 4894 |
| Db | 781 | ATTTAGATGAAATTGTATAAATTGTGTGTTTTTATGGTTTTAATTAAATT | 840 |
| Qy | 4895 | TGAGAAATTAAATTCTGCAACTTGGACAGTCATGCTCTGAATATCTGTACT | 4954 |
| Db | 841 | TGAGAAATTAAATTGTGTAATTAGGATAAGTTGTGAAATTGTGACT | 900 |
| Qy | 4955 | TGAGGAGAAATTATGTTATATTGCCAAATAAAATAAGTTGCAAGTTTTCTTG | 5014 |
| Db | 901 | TGAGGAGAAATTGTGTAATTGTGAAATTGTGAAATTGTGAAATTGTGACT | 960 |
| Qy | 5015 | CCCCAAAGAGCTCTGTCCTTGAACTAAATAACCGCTATGCTGTTAAATTAT | 5074 |
| Db | 961 | TTTAAAGAGTTTGTTGAAATTAAATAATTAATTAATTAATTAATTAATTAAT | 1020 |
| Qy | 5075 | TGGCAAATGCCATTTCACCTAAGGAAATCCATAAGTAACGATACACAA | 5134 |
| Db | 1021 | TGGAAATGTTTAAATTAAAGGAAATTATAAGTAATGATATAATTAATAAA | 1080 |
| Qy | 5135 | AGGTTACTGTTAACGGCATGGCTGAAGATAAAAGAATTTCGCAATGATTTC | 5194 |
| Db | 1081 | AGGTTACTGTTAACGGCATGGTTAAAGGATATAAAAGAATTTCGCAATGATTTC | 1140 |
| Qy | 5195 | CATATGTTGCTCCACCACTGCCAAATAACA | 5224 |
| Db | 1141 | TATATGTTTATTATGTTAAAGGATATAAAAGAATTTCGCAATGATTTC | 1170 |

Search completed: August 7, 2005, 08:16:13
 Job time : 3144.83 sec

GenCore version 5.1.6
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OM nucleic - nucleic search, using SW model

Run on: August 6, 2005, 16:35:01 ; Search time 808.744 Seconds
(without alignments) 10569.369 Million cell updates/sec

Title: US-09-509-591-2
Perfect score: 5224
Sequence: 1 GAATTCCTAGAAATGCGG.....TTCCACCACTGCCATRACA 5224

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 1202784 seqB, 818138359 residues

Total number of hits satisfying chosen parameters: 2405568

Minimum DB seq length: 0

Maximum DB seq length: 200000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Issued Patents NA:*

- 1: /cgns_6/ptodata/1/ina/5A_COMB.seq: *
- 2: /cgns_6/ptodata/1/ina/5B_COMB.seq: *
- 3: /cgns_6/ptodata/1/ina/6A_COMB.seq: *
- 4: /cgns_6/ptodata/1/ina/6B_COMB.seq: *
- 5: /cgns_6/ptodata/1/ina/PCITS_COMB.seq: *
- 6: /cgns_6/ptodata/1/ina/bacfileseq: *

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

| Result No. | Score | Query Match Length | DB ID | Description |
|------------|--------|--------------------|-------|---|
| 1 | 5224 | 100.0 | 5224 | 3 US-09-013-428-2 Sequence 2, Appli |
| 2 | 5224 | 100.0 | 5224 | 3 US-09-013-428-5 Sequence 5, Appli |
| 3 | 5224 | 100.0 | 5224 | 4 US-09-474-699-15 Sequence 15, Appli |
| 4 | 5224 | 100.0 | 5224 | 4 US-09-898-833-2 Sequence 2, Appli |
| 5 | 5224 | 100.0 | 5224 | 4 US-09-151-376-5 Sequence 5, Appli |
| 6 | 5224 | 100.0 | 5224 | 4 US-09-151-376-45 Sequence 45, Appli |
| 7 | 5224 | 100.0 | 5224 | 4 US-09-814-331-16 Sequence 16, Appli |
| 8 | 5224 | 100.0 | 5224 | 1 US-09-148-018A-12 Sequence 12, Appli |
| 9 | 2240 | 42.9 | 2240 | 1 US-08-478-042-12 Sequence 12, Appli |
| 10 | 2240 | 42.9 | 2240 | 1 US-08-445-215-12 Sequence 12, Appli |
| 11 | 2240 | 42.9 | 2240 | 2 US-08-466-612-2 Sequence 12, Appli |
| 12 | 2028.8 | 38.8 | 23497 | 4 US-09-949-056-17072 Sequence 17072, A |
| 13 | 619 | 11.8 | 822 | 3 US-09-033-428-1 Sequence 1, Appli |
| 14 | 619 | 11.8 | 822 | 3 US-09-033-526-6 Sequence 12, Appli |
| 15 | 619 | 11.8 | 822 | 4 US-09-474-699-14 Sequence 14, Appli |
| 16 | 619 | 11.8 | 822 | 4 US-09-698-893-1 Sequence 1, Appli |
| 17 | 619 | 11.8 | 822 | 4 US-09-151-376-6 Sequence 6, Appli |
| 18 | 619 | 11.8 | 822 | 4 US-09-151-376-44 Sequence 44, Appli |
| 19 | 619 | 11.8 | 822 | 4 US-09-814-331-8 Sequence 6, Appli |
| 20 | 409 | 7.8 | 409 | 1 US-08-148-031-8 Sequence 6, Appli |
| 21 | 409 | 7.8 | 409 | 1 US-08-478-042-6 Sequence 6, Appli |
| 22 | 409 | 7.8 | 409 | 1 US-08-645-215-6 Sequence 6, Appli |
| 23 | 409 | 7.8 | 409 | 2 US-08-646-634-6 Sequence 6, Appli |
| 24 | 232.4 | 4.4 | 234 | 1 US-08-148-038A-7 Sequence 7, Appli |
| 25 | 232.4 | 4.4 | 234 | 1 US-08-478-042-7 Sequence 7, Appli |
| 26 | 232.4 | 4.4 | 234 | 1 US-08-645-215-7 Sequence 7, Appli |
| 27 | 232.4 | 4.4 | 234 | 2 US-08-466-604-7 Sequence 7, Appli |

| | | |
|----------|---|-----------------------------------|
| RESULT 1 | US-09-033-428-2 | Sequence 195, Appli |
| | ; Sequence 2, Application US/09033428 | Sequence 14, Appli |
| | ; Patent No. 6254862 | Sequence 3, Appli |
| | ; GENERAL INFORMATION: | Sequence 3, Appli |
| | ; APPLICANT: Lamparski, Henry | Sequence 15371, A |
| | ; APPLICANT: Schinur, Eric | Sequence 14578, A |
| | ; ATTORNEY: Henderson, Daniel | Sequence 14624, A |
| | ; TITLE OF INVENTION: ADENOVIRUS VECTORS SPECIFIC FOR CELLS EXPRESSING ALPHA-FETOPROTEIN AND METHODS OF USE THEREOF | Sequence 15836, A |
| | ; NUMBER OF SEQUENCES: 23 | Sequence 15371, A |
| | ; CORRESPONDENCE ADDRESS: | Sequence 15372, A |
| | ; ADDRESSEE: MORRISON & FOERSTER | Sequence 99, Appli |
| | ; STREET: 755 PAGE MILL ROAD | Sequence 196355, Sequence 196366, |
| | ; CITY: PALO ALTO | |
| | ; STATE: CA | |
| | ; COUNTRY: USA | |
| | ; ZIP: 94304-1018 | |
| | ; COMPUTER READABLE FORM: | |
| | ; MEDIUM TYPE: Floppy disk | |
| | ; COMPUTER: IBM PC compatible | |
| | ; OPERATING SYSTEM: PC-DOS/MS-DOS | |
| | ; SOFTWARE: PatentIn Release #1.0, Version #1.3.0 | |
| | ; CURRENT APPLICATION DATA: | |
| | ; APPLICATION NUMBER: US/09/033, 428 | |
| | ; FILING DATE: | |
| | ; CLASSIFICATION: | |
| | ; ATTORNEY/AGENT INFORMATION: | |
| | ; NAME: POLIZZI, CATHERINE M. | |
| | ; REGISTRATION NUMBER: 40,130 | |
| | ; REFERENCE/DOCKET NUMBER: 34802-30004.00 | |
| | ; TELECOMMUNICATION INFORMATION: | |
| | ; TELEPHONE: (415) 813-5600 | |
| | ; TELEFAX: (415) 494-0792 | |
| | ; TELEX: 706141 MIRENFORIS SFO | |
| | ; INFORMATION FOR SEQ ID NO: 2: | |
| | ; SEQUENCE CHARACTERISTICS: | |
| | ; LENGTH: 5224 base pairs | |
| | ; TYPE: nucleic acid | |
| | ; STRANDEDNESS: single | |
| | ; TOPOLOGY: linear | |
| | ; US-09-033-428-2 | |
| QV | 1 GAATTCCTAGAAATGGGCTAGGGTGTGGTAACTGTTACCCATAGGT | 60 |

Db 1 GATATTCTTAGAAATATGGGTTAGGGTGGTGTGTTACCCCATGGT 60
 Qy 61 GAGATAGCATGGGTAATGCTTCACACACATCATTCAAGATTAAG 120
 Db 61 GAGATAGCATGGTAACTGCTTCACACACATCATTCAAGATTAAG 120
 Qy 121 AACAGACTATGGCTGGAGGTTAATGCTTCACACACATCATTCAAGATTAAG 120
 Db 121 AACAGACTATGGCTGGAGGTTAATGCTTCACACACATCATTCAAGATTAAG 180
 Qy 181 GTTCTATGGGCTGGTTAACGCTGGCAACTTGCAACAGCTGGCTGCC 240
 Db 181 GTTCTATGGGCTGGTTAACGCTGGCAACTTGCAACAGCTGGCTGCC 240
 Qy 241 AAGCCAAAGTACTGCTCTTCAATCGTTGGGCTCTGGGTGATACT 300
 Db 241 AAGCCAAAGTACTGCTCTTCAATCGTTGGGCTCTGGGTGATACT 300
 Qy 301 GAGAAATATACATTCATATGCTGGAGATGAGATGAGATGAGATG 360
 Db 301 GAGAAATATACATTCATATGCTGGAGATGAGATGAGATGAGATG 360
 Qy 361 TCATTGATATGATGATGAGGACAATAGTGTATAATCCTTAGTACACAT 420
 Db 361 TCATTGATATGATGATGAGGACAATAGTGTATAATCCTTAGTACACAT 420
 Qy 421 GAGGGAGGGTGGTACTTCAATTCTTAACTGTTCTGGTGAATCT 480
 Db 421 GAGGGAGGGTGGTACTTCAATTCTTAACTGTTCTGGTGAATCT 480
 Qy 481 TCTCTGACAAATGTTGGGGACACAGGAACTCACAGATGAGCTTGATT 540
 Db 481 TCTCTGACAAATGTTGGGGACACAGGAACTCACAGATGAGCTTGATT 540
 Qy 541 AGAAAGTAAATGGCCAGGAATAATACAGCTAACATGACTGATACCAA 600
 Db 541 AGAAAGTAAATGGCCAGGAATAATACAGCTAACATGACTGATACCAA 600
 Qy 601 ACAAGGGTTCATGAAATACTGAAGGTTAATAGTGTGCAAGGGAGGG 660
 Db 601 ACAAGGGTTCATGAAATACTGAAGGTTAATAGTGTGCAAGGGAGGG 660
 Qy 661 CTCTCTCTACTAGACATACATCAGGTTAATGGTTAATAGTGTGCAAGGG 720
 Db 661 CTCTCTCTACTAGACATACATCAGGTTAATGGTTAATAGTGTGCAAGGG 720
 Qy 721 TAAGGGTGGTACATTGTCAGCATGAGGAGGAGGAGGAGGAGGAGG 780
 Db 721 TAAGGGTGGTACATTGTCAGCATGAGGAGGAGGAGGAGGAGGAGG 780
 Qy 781 ATTTCACATACCAACCACCAATCACAAACAAAATGAAAGAATCTCAGAAC 840
 Db 781 ATTTCACATACCAACCACCAATCACAAACAAAATGAAAGAATCTCAGAAC 840
 Qy 841 GTGAGATAGGAGAATTCTCACAAACCAACGAGTCACTGCTGGAGAA 900
 Db 841 GTGAGATAGGAGAATTCTCACAAACCAACGAGTCACTGCTGGAGAA 900
 Qy 901 GTATATCATATATTPACCTAACATCATGCTTAATGATAATATCTGATT 960
 Db 901 GTATATCATATTPACCTAACATCATGCTTAATGATAATATCTGATT 960
 Qy 961 TTAATGCTATAGTACCGCATTAGAAGATATTCTCATTTATTCACAAATA 1020
 Db 961 TTAATGCTATAGTACCGCATTAGAAGATATTCTCATTTATTCACAAATA 1020
 Qy 1021 ACTTGGGGTATAGCATTTCTCATGATATGAGAAATTAACACAGTTGATT 1080
 Db 1021 ACTTGGGGTATAGCATTTCTCATGATATGAGAAATTAACACAGTTGATT 1080
 Qy 1081 TTGCCCTGTCATACAGTAAATGACCATAGACATTAGATTAAATGTTGAT 1140
 Db 1081 TTGCCCTGTCATACAGTAAATGACCATAGACATTAGATTAAATGTTGAT 1140

Qy 1141 CTTCTATACCAAGTCAGTTACTGTCAGTTGCTCTGAGGCTCACAGCT 1200
 Db 1141 CTTCTATACCAAGTCAGTTACTGTCAGTTGCTCTGAGGCTCACAGCT 1200
 Qy 1201 TATGAAATACGAACTGAGATTACATCAATGCAAAAGCTGGCTGAACTGT 1260
 Db 1201 TATGAAATACGAACTGAGATTACATCAATGCAAAAGCTGGCTGAACTGT 1260
 Qy 1261 ACTTAGGACTAACTTGAGGAACTACACATAGTGGATGTTGCTGTACAT 1320
 Db 1261 ACTTAGGACTAACTTGAGGAACTACACATAGTGGATGTTGCTGTACAT 1320
 Qy 1321 ACAACTCTGGTCAAGCTCAGATTCATGTCAGTCAAAAGCTGGCTGAACTGT 1380
 Db 1321 ACAACTCTGGTCAAGCTCAGATTCATGTCAGTCAAAAGCTGGCTGAACTGT 1380
 Qy 1381 GGTTCCTTCACTGTCATCTATTTCTCAACCACTCACAGTGCTGAACTGT 1440
 Db 1381 GGTTCTCTTCACTGTCATCTATTTCTCAACCACTCACAGTGCTGAACTGT 1440
 Qy 1441 TGCAGCTTGTGATCCAAATCTGCTCTGCTCTGCTGCTGCTGCTGCTGCT 1500
 Db 1441 TGCAGCTTGTGATCCAAATCTGCTCTGCTCTGCTGCTGCTGCTGCTGCT 1500
 Qy 1501 AGTAGTAACTGATCCAAATCTGCTCTGCTCTGCTGCTGCTGCTGCTGCT 1560
 Db 1501 AGTAGTAACTGATCCAAATCTGCTCTGCTCTGCTGCTGCTGCTGCTGCT 1560
 Qy 1561 CAACGGGGAGTTGCGGAATGTTCTCAAATGTCAGTCACTGCTGCTGCTGCT 1620
 Db 1561 CAACGGGGAGTTGCGGAATGTTCTCAAATGTCAGTCACTGCTGCTGCTGCT 1620
 Qy 1621 TCTGTTCTTAAATCTTAATCTTAACTTCTGCTGCTGCTGCTGCTGCTGCT 1680
 Db 1621 TCTGTTCTTAAATCTTAATCTTAACTTCTGCTGCTGCTGCTGCTGCTGCT 1680
 Qy 1680 1680
 Db 1680 1680
 Qy 1681 AGAGTGAAGGGAGCTGTTAATACAGCTAACAGGATAGGAGGAGG 1740
 Db 1681 AGAGTGAAGGGAGCTGTTAATACAGCTAACAGGATAGGAGGAGG 1740
 Qy 1741 GTTGGGTTAACTGGTCACTTATCTTAATCTAACAAACTGACAGTCACTT 1800
 Db 1741 GTTGGGTTAACTGGTCACTTATCTTAATCTAACAAACTGACAGTCACTT 1800
 Qy 1801 AGTACTAAGCTTGCATTATCTCATACACTAACATAATTCACACGACTT 1860
 Db 1801 AGTACTAAGCTTGCATTATCTCATACACTAACATAATTCACACGACTT 1860
 Qy 1860 1860
 Db 1860 1860
 Qy 1861 TTGACAGTATTGGAAACTCTCTACTGTCCTTATCATAGCTTATCCCTT 1920
 Db 1861 TTGACAGTATTGGAAACTCTCTACTGTCCTTATCATAGCTTATCCCTT 1920
 Qy 1921 TGAACACAAAGAGACAGTTCAAAATAAAATGATTTTATAGTCCTTGT 1980
 Db 1921 TGAACACAAAGAGACAGTTCAAAATAAAATGATTTTATAGTCCTTGT 1980
 Qy 1980 1980
 Db 1980 1980
 Qy 1981 CTATATAGTCCAGAAGGGTTAACCTAACATGATTATAGCTCCCTTGT 2040
 Db 1981 CTATATAGTCCAGAAGGGTTAACCTAACATGATTATAGCTCCCTTGT 2040
 Qy 2040 2040
 Db 2040 2040
 Qy 2041 TGCCACTTGCAGGAACTCCAAATCTAGTATTCTACATTAACCTTGCCCT 2100
 Db 2041 TGCCACTTGCAGGAACTCCAAATCTAGTATTCTACATTAACCTTGCCCT 2100
 Qy 2101 TTCAAAATCTGCTTCTCATCCCTAGTGTGCTATGTCCTTACCGGTTGCT 2160
 Db 2101 TTCAAAATCTGCTTCTCATCCCTAGTGTGCTATGTCCTTACCGGTTGCT 2160
 Qy 2160 2160
 Db 2160 2160
 Qy 2161 TTCCACACCTTCTCATCCCTAGGACACTAACCTCCCTTATTCGGCCAC 2220
 Db 2161 TTCCACACCTTCTCATCCCTAGGACACTAACCTCCCTTATTCGGCCAC 2220

CITY: Palo Alto
 STATE: CA
 COUNTRY: USA
 Z.I.P: 94304-1018

COMPUTER READABLE FORM:
 COMPUTER: IBM Compatible
 OPERATING SYSTEM: Windows
 SOFTWARE: FastSeq for Windows Version 2.0b

CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/09/033, 556
 FILING DATE:
 CLASSIFICATION:
 PRIORITY APPLICATION DATA:
 APPLICATION NUMBER:
 FILING DATE:
 ATTORNEY/AGENT INFORMATION:
 NAME: Catherine Polizzi M
 REGISTRATION NUMBER: 40, 30
 REFERENCE/DOCKET NUMBER: 34802-2010.00
 TELECOMMUNICATION INFORMATION:
 TELEPHONE: 650-813-5600
 TELEFAX: 650-494-0792

INFORMATION FOR SEQ ID NO: 5:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 5224 base pairs
 TYPE: nucleic acid
 STRANDEDNESS: single
 TOPOLOGY: linear

US-09-033-556-5

Query Match 100.0%; Score 5224; DB 3; Length 5224;
 Best Local Similarity 100.0%; Pred. No. 0;
 Matches 5224; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 GAATCTTGAATATGCGCTTAGGGTGTGGTAACTTCACCCATACT 60
 DB 1 GAATTCTTGAATATGCGCTTAGGGTGTGGTAACTTCACCCATACT 60
 QY 61 GAGATAGCAATTGGTTAAATGGCTTCAACACACATTCATAAGAATTAGG 120
 DB 61 GAGATAGCAATTGGTTAAATGGCTTCAACACACATTCATAAGAATTAGG 120
 QY 121 AACAGACTATGGCTGGAGACTTGAGGATGTGCTGTCATAACACTGGGTATCT 180
 DB 121 AACAGACTATGGCTGGAGACTTGAGGATGTGCTGTCATAACACTGGGTATCT 180
 QY 181 GTTCATGGCTGGAGACTTGAGGATGTGCTGTCATAACACTGGGTATCT 240
 DB 181 GTTCATGGCTGGAGACTTGAGGATGTGCTGTCATAACACTGGGTATCT 240
 QY 241 AAGCCAGGACTACTGCCTCTTCAATCAGTTGTTGGGCCCTGGGGCTGATACT 300
 DB 241 AAGCCAGGACTACTGCCTCTTCAATCAGTTGTTGGGCCCTGGGGCTGATACT 300
 QY 301 GAGAAATAAACCTTCAATGTCCTGGTGGAGATGAGATGAGATGAGATGTCAT 360
 DB 301 GAGAAATAAACCTTCAATGTCCTGGTGGAGATGAGATGAGATGAGATGTCAT 360
 QY 361 TCATTGTCATGAACTGATGAGGACATTAGTAACTAACCTTGTGACACCT 420
 DB 361 TCATTGTCATGAACTGATGAGGACATTAGTAACTAACCTTGTGACACCT 420
 QY 421 GAGGGTAGGGTGTGACTATCAATTCTTCAATGTCCTGGTGGAGATGAGATGAGATGTCAT 480
 DB 421 GAGGGTAGGGTGTGACTATCAATTCTTCAATGTCCTGGTGGAGATGAGATGAGATGTCAT 480
 QY 481 TGCTGTGACAATGTTGTCGGACCACAGGATCACAATGAGCTTGAATT 540
 DB 481 TGCTGTGACAATGTTGTCGGACCACAGGATCACAATGAGCTTGAATT 540
 QY 541 AAGAAGTTAACCTTCAATGTCCTGGTGGAGATGAGATGAGATGTCAT 600

RESULT 2

US-09-033-556-5

Sequence 5, Application US/09033556

Patent No. 6432700

GENERAL INFORMATION:

APPLICANT: Henderson, Daniel R.

APPLICANT: Yu, De Chao

TITLE OF INVENTION: ADENOVIRUS VECTORS CONTAINING HETEROLOGOUS TRANSCRIPTION REGULATORY ELEMENTS AND METHODS

TITLE OF INVENTION: HETEROLOGOUS TRANSCRIPTION REGULATORY ELEMENTS AND METHODS

NUMBER OF SEQUENCES: 41

CORRESPONDENCE ADDRESS:

ADRESSE: MORRISON & FOERSTER

STREET: 755 PAGE MILL ROAD

| | | | |
|------|--|------|----|
| 541 | AAGAGGTTATGGTCAGGATAATTACATAGCTAACAGACTATACATCAA | 600 | Qy |
| 601 | ACAAGAGGTTCCATGAGAAATACTGAAGGTTAATAGTTCAGGTGAGGG | 660 | Db |
| 601 | ACAAGAGGTTCCATGAGAAATACTGAAGGTTAATAGTTCAGGTGAGGG | 660 | Qy |
| 661 | CTCTCTCTGACTAGAGACTATCGAAATACATCAGGAAATATTGAGGT | 720 | Db |
| 661 | CTCTCTCTGACTAGAGACTATCGAAATACATCAGGAAATATTGAGGT | 720 | Qy |
| 721 | TAAGGGTGGTACTTTGTCAGCATGCATGAGAGAGTGAACTTGGAAAC | 780 | Db |
| 721 | TAAGGGTGGTACTTTGTCAGCATGCATGAGAGAGTGAACTTGGAAAC | 780 | Qy |
| 781 | ATTTCACTAACACCACCCATTCAAACAAACAAATGAGAACTCAGAAC | 840 | Db |
| 781 | ATTTCACTAACACCACCCATTCAAACAAACAAATGAGAACTCAGAAC | 840 | Qy |
| 841 | GTGAGATAAGAGAGGAATTTCACACCCACCGTAGCTAACGCTGAGA | 900 | Db |
| 841 | GTGAGATAAGAGAGGAATTTCACACCCACCGTAGCTAACGCTGAGA | 900 | Qy |
| 901 | GTATPATCPATAATTACACTACATCATGATAATGATAATGATAAT | 960 | Db |
| 901 | GTATPATCPATAATTACACTACATCATGATAATGATAATGATAAT | 960 | Qy |
| 961 | TTTAACTGCTATAAGACCCAGGTTAGAGATAATTCCATTATTCAGAA | 1020 | Db |
| 961 | TTTAACTGCTATAAGACCCAGGTTAGAGATAATTCCATTATTCAGAA | 1020 | Qy |
| 1021 | ACTTAGGGGATGATCATGCTAATTCTGATGATATGAGAAATTAAAC | 1080 | Db |
| 1021 | ACTTAGGGGATGATCATGCTAATTCTGATGATATGAGAAATTAAAC | 1080 | Qy |
| 1081 | TTCCTGTCATACAGCTAATTGACCTAAGGAAATTAGATGATTGAT | 1140 | Db |
| 1081 | TTCCTGTCATACAGCTAATTGACCTAAGGAAATTAGATGATTGAT | 1140 | Qy |
| 1141 | CTTCTTAATACCAAGTTCAGTTAGTTCAGTGTCCATGAGCT | 1200 | Db |
| 1141 | CTTCTTAATACCAAGTTCAGTTAGTTCAGTGTCCATGAGCT | 1200 | Qy |
| 1201 | TATGAAAGTAACGGAACTGAAATTACATCACTGAAAGCTGTGAACT | 1260 | Db |
| 1201 | TATGAAAGTAACGGAACTGAAATTACATCACTGAAAGCTGTGAACT | 1260 | Qy |
| 1261 | ACTTAGGACTAACTTGACCAATAACACATGATTGAGGTGTTGCTG | 1320 | Db |
| 1261 | ACTTAGGACTAACTTGACCAATAACACATGATTGAGGTGTTGCTG | 1320 | Qy |
| 1321 | ACAACCTCTGGTCAAGCTCTTTATGCTGTCAGAAATTGCTGTT | 1380 | Db |
| 1321 | ACAACCTCTGGTCAAGCTCTTTATGCTGTCAGAAATTGCTGTT | 1380 | Qy |
| 1381 | GGTTCTCTTCACTGCTATTTCTCACCACTCACATGCTACAACTG | 1440 | Db |
| 1381 | GGTTCTCTTCACTGCTATTTCTCACCACTCACATGCTACAACTG | 1440 | Qy |
| 1441 | TGCAAGCTATGATCCAAATACTATCTGCTCACTGTGTCAGAAAG | 1500 | Db |
| 1441 | TGCAAGCTATGATCCAAATACTATCTGCTCACTGTGTCAGAAAG | 1500 | Qy |
| 1501 | AGTAGTATCAAAAGCACATCAACGCTCCACTGGAGGCTTAAGAG | 1560 | Db |
| 1501 | AGTAGTATCAAAAGCACATCAACGCTCCACTGGAGGCTTAAGAG | 1560 | Qy |
| 1561 | CAAACTGGGAGTTTGCTGGAAATGTCCTAAATGTCCTGGAGGTC | 1620 | Db |
| 1561 | CAAACTGGGAGTTTGCTGGAGTTTCCTAAATGTCCTGGAGGTC | 1620 | Qy |
| 1621 | TCTGTCTCTAAACTTAACTTAACTTAACTGCCCCGTCATCCCAC | 1680 | Db |
| 1621 | TCTGTCTCTAAACTTAACTTAACTTAACTGCCCCGTCATCCCAC | 1680 | Qy |
| 1681 | AGAGGAAAGGGCCGATTAATTAACACTAGCTAAGCTAGGCCAG | 1740 | Db |
| 1681 | AGAGGAAAGGGCCGATTAATTAACACTAGCTAAGCTAGGCCAG | 1740 | Qy |
| 1741 | GTTGGTAACCTGGTCACTTATCTTAACAAATTCAGGCTTACAT | 1800 | Db |
| 1741 | GTTGGTAACCTGGTCACTTATCTTAACAAATTCAGGCTTACAT | 1800 | Qy |
| 1801 | AGTTACTAAGCTCTGACTTATCTCATACCTACACTCGCTTAT | 1860 | Db |
| 1801 | AGTTACTAAGCTCTGACTTATCTCATACCTACACTCGCTTAT | 1860 | Qy |
| 1861 | TGACAGTATTGCGAAACTCTCAACTGGCTCCTTATCATAGCTT | 1920 | Db |
| 1861 | TGACAGTATTGCGAAACTCTCAACTGGCTCCTTATCATAGCTT | 1920 | Qy |
| 1921 | TGAAACAAAGAGACAGTTCAAAATCACAAATGATGAACTCAG | 1980 | Db |
| 1921 | TGAAACAAAGAGACAGTTCAAAATCACAAATGATGAACTCAG | 1980 | Qy |
| 1981 | CTATPATAGTCCAGAGGTTAAACTCCATTAAAGTGTGTTAG | 2040 | Db |
| 1981 | CTATPATAGTCCAGAGGTTAAACTCCATTAAAGTGTGTTAG | 2040 | Qy |
| 2041 | TGCCACTTGGCAGGAACTCCATTCTAGTATTCTCTATTAAC | 2100 | Db |
| 2041 | TGCCACTTGGCAGGAACTCCATTCTAGTATTCTCTATTAAC | 2100 | Qy |
| 2101 | TCAAACACTGCTCTCAGGTTAAACTCCATTAAAGTGTGTTAG | 2160 | Db |
| 2101 | TCAAACACTGCTCTCAGGTTAAACTCCATTAAAGTGTGTTAG | 2160 | Qy |
| 2161 | TTCACACCCTTTACATTCTGACACTATACCTCCCTCTTCA | 2220 | Db |
| 2161 | TTCACACCCTTTACATTCTGACACTATACCTCCCTCTTCA | 2220 | Qy |
| 2221 | CTTAAATTCTCTCAGATCTCATGAGTCTCTCAGGAACTCT | 2280 | Db |
| 2221 | CTTAAATTCTCTCAGATCTCATGAGTCTCTCAGGAACTCT | 2280 | Qy |
| 2281 | CCTCCAAGATGTCATGACTTCTCTCTCTCTCTCTCTCT | 2340 | Db |
| 2281 | CCTCCAAGATGTCATGACTTCTCTCTCTCTCTCTCTCT | 2340 | Qy |
| 2341 | TGTTGTAATCATCATGATCTGTTCTGTTCTGTTCTGTTCT | 2400 | Db |
| 2341 | TGTTGTAATCATCATGATCTGTTCTGTTCTGTTCTGTTCT | 2400 | Qy |
| 2401 | TACATGGTCCCTCTCTCTGTTCTGTTCTGTTCTGTTCT | 2460 | Db |
| 2401 | TACATGGTCCCTCTCTGTTCTGTTCTGTTCTGTTCTGTTCT | 2460 | Qy |
| 2461 | GACTTACATTATGATGATAATAAACCCATCTATGAGCTAT | 2520 | Db |
| 2461 | GACTTACATTATGATGATAATAAACCCATCTATGAGCTAT | 2520 | Qy |
| 2521 | AGACCCGGTTCTGGCAATTATGTTATTTATCTCTTACCAT | 2580 | Db |
| 2521 | AGACCCGGTTCTGGCAATTATGTTATTTATCTCTTACCAT | 2580 | Qy |
| 2581 | GAGGACTACTATCTTATGTTGATAAGATAAGCCAGAGATG | 2640 | Db |
| 2581 | GAGGACTACTATCTTATGTTGATAAGATAAGCCAGAGATG | 2640 | Qy |
| 2641 | ACTCACCAAGTCATGTTGAGCAAGGCAAAATCAACCGTCTCC | 2700 | Db |
| 2641 | ACTCACCAAGTCATGTTGAGCAAGGCAAAATCAACCGTCTCC | 2700 | Qy |
| 2701 | TTAGGTGTTAACTGCTACTGCTACTGCTACTGCTACTGCT | 2760 | Db |
| 2701 | TTAGGTGTTAACTGCTACTGCTACTGCTACTGCTACTGCT | 2760 | Qy |


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Db 4921 TAGGCCAAGCTCATGCTTGAATTCGTGTTGAGGAAATTGGTATATTGC 4980 Qy
Qy 4981 AAATAATAATAGTTGCAAGTTTCTGCCAAAGAGCTGTCGTCCTGAC 5040 Db
Db 4981 AAATAATAATAGTTGCAAGTTTCTGCCAAAGAGCTGTCGTCCTGAC 5040 Qy
Qy 5041 ATAAATAACATAACCGCTATGCTTAATTGGCAATGTCCTTCACTAA 5100 Db
Db 5041 ATAAATAACATAACCGCTATGCTTAATTGGCAATGTCCTTCACTAA 5100 Qy
Qy 5101 GGAAATACATAAACGCTATGCTTAATTGGCAATGTCCTTCACTAA 5100 Db
Db 5101 GGAAATACATAAACGCTATGCTTAATTGGCAATGTCCTTCACTAA 5100 Qy
Qy 5161 GAAAGAGATAAAGAATTCAAGATTAACAGATAACCAACAGGTTACTAGTAACAGGCAATGCT 5160 Db
Db 5161 GAAAGAGATAAAGAATTCAAGATTAACAGATAACCAACAGGTTACTAGTAACAGGCAATGCT 5160 Qy
Qy 5221 AACA 5224 Db
Db 5221 AACA 5224 Qy
Db 5221 AACA 5224 Qy
Db ; RESULT 3
US-09-474-699-15
; Sequence 15, Application US/09474699
; Patent NO. 6495130
; GENERAL INFORMATION:
; APPLICANT: Henderson, Daniel R.
; APPLICANT: Yu, De Chao
; TITLE OF INVENTION: TARGET CELL-SPECIFIC ADENOVIRAL VECTORS
; TITLE OF INVENTION: CONTAINING E3 AND METHODS OF USE THEREOF
; FILE REFERENCE: 348022001300
; CURRENT APPLICATION NUMBER: US/09/474,699
; CURRENT FILING DATE: 1999-12-29
; PRIORITY APPLICATION NUMBER: 60/114,262
; PRIORITY FILING DATE: 1998-12-30
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: FASTSEQ for Windows Version 4.0
; SEQ ID NO 15
; LENGTH: 5224
; TYPE: DNA
; ORGANISM: Homo Sapien
; US-09-474-699-15

Query Match 100.0%; Score 5224; DB 4; Length 5224;
Best Local Similarity 100.0%; Prod. No. 0; Mismatches 0; Indels 0; Gaps 0;
Matches 5224; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 GAACTTCTGAAATAGGGGTAGGGGTGGTGTGTTAACCTCCATAGGT 60 Db
Db 1 GAACTTCTGAAATAGGGGTAGGGGTGGTGTGTTAACCTCCATAGGT 60 Qy
Qy 61 GAGATTAAGCATGGTTAAATGTCCTTCACACACACATTCATAGAATTAAGC 120 Db
Db 61 GAGATTAAGCATGGTTAAATGTCCTTCACACACACATTCATAGAATTAAGC 120 Qy
Qy 121 AACGACTATGGCTGGAGACTTGAGATGCTCTACACTGGGTATCT 180 Db
Db 121 AACGACTATGGCTGGAGACTTGAGATGCTCTACACTGGGTATCT 180 Qy
Qy 181 GTCTCTATGGGCTGTGTTAACCTGAACTGCAAGGGTCACTGACTTCCCC 240 Db
Db 181 GTCTCTATGGGCTGTGTTAACCTGAACTGCAAGGGTCACTGACTTCCCC 240 Qy
Qy 241 AACCCCAAGGACTCTGCTCTTCTATCTGTTGGGGCTCTGGGGCTGTGATATCT 300 Db
Db 241 AACCCCAAGGACTCTGCTCTTCTATCTGTTGGGGCTCTGGGGCTGTGATATCT 300 Qy
Qy 301 GAGAAATAACATTCATAGTCGTTGAGATGAGATGAGATGAGATGTCAT 360 Db
Db 301 GAGAAATAACATTCATAGTCGTTGAGATGAGATGAGATGTCAT 360 Qy
Db ; 361 TCAATTGATATGATGATGAGGAAATTGGTATACTCTAGTACACAACT 420
Db 361 TCAATTGATATGATGATGAGGAAATTGGTATACTCTAGTACACAACT 420 Qy
Db 421 GAGGTAGGGTGTCTATCAATTCTATTAAAGAATGACTTCTATTATTA 480
Db 421 GAGGTAGGGTGTCTATCAATTCTATTAAAGAATGACTTCTATTATTA 480 Qy
Qy 481 TCTCTGACAAATGTTGTTGGACACAGGATCACAAAGATGAGCTTGATT 540
Db 481 TCTCTGACAAATGTTGTTGGACACAGGATCACAAAGATGAGCTTGATT 540 Qy
Qy 541 AGAAGTTAATGTCAGGAAATTACATGCTTACAAATGACTATGATACATCAA 600
Db 541 AGAAGTTAATGTCAGGAAATTACATGCTTACAAATGACTATGATACATCAA 600 Qy
Qy 601 ACAAGAGGTCCATGAAATAATCTGAAAGGTTATAGTTGTCAAAGGTGAGGG 660
Db 601 ACAAGAGGTCCATGAAATAATCTGAAAGGTTATAGTTGTCAAAGGTGAGGG 660 Qy
Qy 661 CTCTCTCTACTAGTAGAGACTATCAGAAATCTAGGATATTATGATAGCT 720
Db 661 CTCTCTCTACTAGTAGAGACTATCAGAAATCTAGGATATTATGATAGCT 720 Qy
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Db 721 TAAGGTTGCTGACATTTGTCACGATTGTTGAGGAGGAGTATTGAAAC 780 Qy
Qy 781 ATTTCACAACTACACCACCCAACTCAACAAACAAAAAGGAAATTCAGAAC 840
Db 781 ATTTCACAACTACACCACCCAACTCAACAAACAAAAAGGAAATTCAGAAC 840 Qy
Qy 841 GTGAGATAAGGAGATTCTCACACCCAACTCAACAAACAAAAAGGAAATTCAGAAC 900
Db 841 GTGAGATAAGGAGATTCTCACACCCAACTCAACAAACAAAAAGGAAATTCAGAAC 900 Qy
Qy 901 GTATATCTAAATTACATGCTTAATGATTAATGATAATATCTGCTGAGAA 960
Db 901 GTATATCTAAATTACATGCTTAATGATTAATGATAATATCTGCTGAGAA 960 Qy
Qy 961 TTAAATGCTTAATGATCACCACCACTCAACAAACAAAAAGGAAATTCAGAAC 1020
Db 961 TTAAATGCTTAATGATCACCACCACTCAACAAACAAAAAGGAAATTCAGAAC 1020 Qy
Qy 1021 ACTTCAGGGGATAGTCATTTCATGATATGAGAAATTAAACAGITGGATT 1080
Db 1021 ACTTCAGGGGATAGTCATTTCATGATATGAGAAATTAAACAGITGGATT 1080 Qy
Qy 1081 TTGCTCTGCAACAGSTAATTGACCATAGACAACTTAAATTGTTGAAAT 1140
Db 1081 TTGCTCTGCAACAGSTAATTGACCATAGACAACTTAAATTGTTGAAAT 1140 Qy
Qy 1141 CTTCTTAATACCAAAAGTTCAGTTACTGTCATGTTGCTCTGAGTGGCTCACAGCT 1200
Db 1141 CTTCTTAATACCAAAAGTTCAGTTACTGTCATGTTGCTCTGAGTGGCTCACAGCT 1200 Qy
Qy 1201 TATGAAAGTACAGGAAATCAAGAAATTACATCAATGCAAGCATGTTGCTGT 1260
Db 1201 TATGAAAGTACAGGAAATCAAGAAATTACATCAATGCAAGCATGTTGCTGT 1260 Qy
Qy 1261 ACTTAGGACTAACTTGACCAACACATAGTGAGATCTTGGTTGACAT 1320
Db 1261 ACTTAGGACTAACTTGACCAACACATAGTGAGATCTTGGTTGACAT 1320 Qy
Qy 1261 ACTTAGGACTAACTTGACCAACACATAGTGAGATCTTGGTTGACAT 1320
Db 1261 ACTTAGGACTAACTTGACCAACACATAGTGAGATCTTGGTTGACAT 1320 Qy
Qy 1321 ACAACTCTGGTCACAGCTCTCTTATGCTGCTCTGGAAATTGGTCTCT 1380
Db 1321 ACAACTCTGGTCACAGCTCTCTTATGCTGCTCTGGAAATTGGTCTCT 1380 Qy
Qy 1321 ACAACTCTGGTCACAGCTCTCTTATGCTGCTCTGGAAATTGGTCTCT 1380
Db 1321 ACAACTCTGGTCACAGCTCTCTTATGCTGCTCTGGAAATTGGTCTCT 1380 Qy
Qy 1381 GTTTCCTCTTCTACTGCTATCTTCTACCACTCACATGCTACATACTGTC 1440
Db 1381 GTTTCCTCTTCTACTGCTATCTTCTACCACTCACATGCTACATACTGTC 1440 Qy
Qy 1381 GTTTCCTCTTCTACTGCTATCTTCTACCACTCACATGCTACATACTGTC 1440

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| | | | | |
|----|----|------|---|------|
| Db | Qy | 3601 | GCCCTGTTTGTAACTAATGTCCTTGAGGATGTTACTGATAGCGAACCTTGT | 3720 |
| Db | Qy | 3661 | AATGGACAAAACCTACAATGATGGGATGTTACTGATAGCGAACCTTGT | 3720 |
| Db | Qy | 3661 | AATGGACAAAACCTACAATGATGGGATGTTACTGATAGCGAACCTTGT | 3720 |
| Db | Qy | 3721 | TATCTATGATAAATGTTATGCTGAGCTACTGACGGTAACCTTGT | 3780 |
| Db | Qy | 3721 | TATCTATGATAAATGTTATGCTGAGCTACTGACGGTAACCTTGT | 3780 |
| Db | Qy | 3781 | TAATGTCGCCCTAGTAGCTGCGATGATCTTTTAAGTACTGACTGCTA | 3840 |
| Db | Qy | 3781 | TAATGTCGCCCTAGTAGCTGCGATGATCTTTTAAGTACTGACTGCTA | 3840 |
| Db | Qy | 3841 | TTTAAATTGTTAAATGCTAGGCCTTCATGAGAAGTTGAGAG | 3900 |
| Db | Qy | 3841 | TTTAAATTGTTAAATGCTAGGCCTTCATGAGAAGTTGAGAG | 3900 |
| Db | Qy | 3901 | AGATGAAATACTACTATCTACCTACCTAGAGAACCAATGTTAAACTTGTG | 3960 |
| Db | Qy | 3901 | AGATGAAATACTACTATCTACCTACCTAGAGAACCAATGTTAAACTTGTG | 3960 |
| Db | Qy | 3961 | TCCATTATTCGCTTTATCACAATTTTTAGGGTGGAGGATACAGGAG | 4020 |
| Db | Qy | 3961 | TCCATTATTCGCTTTATCACAATTTTTAGGGTGGAGGATACAGGAG | 4020 |
| Db | Qy | 3961 | TCCATTATTCGCTTTATCACAATTTTTAGGGTGGAGGATACAGGAG | 4020 |
| Db | Qy | 4021 | GTACATGATACAACTGAGACACTTCCTCATGATGTTGCTGTTTCAGTA | 4080 |
| Db | Qy | 4021 | GTACATGATACAACTGAGACACTTCCTCATGATGTTGCTGTTTCAGTA | 4080 |
| Db | Qy | 4081 | ACATATATGAGCATTTCCATTCAAAATCTTCACAAAGTTTGT | 4140 |
| Db | Qy | 4081 | ACATATATGAGCATTTCCATTCAAAATCTTCACAAAGTTTGT | 4140 |
| Db | Qy | 4141 | GGCTGTATATCACCCACTTTGATGATGACCAATTATTTTCTGGTGTGCTA | 4200 |
| Db | Qy | 4141 | GGCTGTATATCACCCACTTTGATGATGACCAATTATTTTCTGGTGTGCTA | 4200 |
| Db | Qy | 4141 | GGCTGTATATCACCCACTTTGATGATGACCAATTATTTTCTGGTGTGCTA | 4200 |
| Db | Qy | 4201 | TTTGTATTTATACTTACCTTGTGATGAAACCTGTGAGACGTTAGATACT | 4260 |
| Db | Qy | 4201 | TTTGTATTTATACTTACCTTGTGATGAAACCTGTGAGACGTTAGATACT | 4260 |
| Db | Qy | 4261 | GTCCTGGCTCACTCCACAGATTCGTTACTGCTGGTTACAGACTGCA | 4320 |
| Db | Qy | 4261 | GTCCTGGCTCACTCCACAGATTCGTTACTGCTGGTTACAGACTGCA | 4320 |
| Db | Qy | 4321 | TTCGGAAATCAAAGTTCCCCCAGTGATCTATGCTGAGGGACCTTG | 4380 |
| Db | Qy | 4321 | TTCGGAAATCAAAGTTCCCCCAGTGATCTATGCTGAGGGACCTTG | 4380 |
| Db | Qy | 4381 | TAGACAGGATGATGGGGTGGCCACTCTAGCATCATTTAGTATRACAT | 4440 |
| Db | Qy | 4381 | TAGACAGGATGATGGGGTGGCCACTCTAGCATCATTTAGTATRACAT | 4440 |
| Db | Qy | 4441 | CATCTGTAGTGTGCTAAGTGATGATGTCACCTTAAAGACACATGTCGAA | 4500 |
| Db | Qy | 4441 | CATCTGTAGTGTGCTAAGTGATGATGTCACCTTAAAGACACATGTCGAA | 4500 |
| Db | Qy | 4501 | TAATATATATGGACTTGTTTATGGCTGTGCTCTAGTTCTANGTTAGCC | 4560 |
| Db | Qy | 4501 | TAATATATATGGACTTGTTTATGGCTGTGCTCTAGTTCTANGTTAGCC | 4560 |
| Db | Qy | 4561 | ATACATCCATCAATCTTAACTGACCATGATGACGATGAAAG | 4620 |
| Db | Qy | 4561 | ATACATCCATCAATCTTAACTGACCATGATGACGATGAAAG | 4620 |
| Db | Qy | 4621 | TGTTCTGAGTAAACATGACGACATATTCAAGCATGATGAAAG | 4680 |
| Db | Qy | 4621 | TGTTCTGAGTAAACATGACGACATATTCAAGCATGATGAAAG | 4680 |
| Db | Qy | 4681 | ATTATCTGCACTTGGACAGTCATCTCTGAAATTGACTTTGAGATTT | 4740 |
| Db | Qy | 4681 | ATTATCTGCACTTGGACAGTCATCTCTGAAATTGACTTTGAGATTT | 4740 |

RESULT 4

S-09-898-883-2
Sequence 2, Application US/0989888J
Patent No. 6585968

GENERAL INFORMATION:
APPLICANT: Little, Andrew
Iamparaki, Henry

Schuur, Eric
Henderson, Daniel

TITLE OF INVENTION: ADENOVIRUS VECTOR EXPRESSING APHIDIN
NUMBER OF SEQUENCES: 23

CORRESPONDENCE ADDRESS:
ADDRESSEE: MORRISON & FOERSTER
STREET: 755 PAGE MILL ROAD

CITY: PALO ALTO
STATE: CA
Country: USA

COONRICK: USA
ZIP: 94304-1018
COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compat.
OPERATING SYSTEM: PC-DOS

SOFTWARE: PatentIn Release #1.0, Version #1.300
CURRENT APPLICATION DATA:
APPLICATION NUMBER: ITC/09/0000 883

RECEIVED NUMBER: 00/00/0000, 0000
FILING DATE: 02-Jul-2001
CLASSIFICATION: <Unknown>

PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/033,428

FILING DATE: <Unknown>
ATTORNEY/AGENT INFORMATION:

NAME: POLIZZI, CATHERINE M.
REGISTRATION NUMBER: 40,130

TELECOMMUNICATION INFORMATION:
 TELEPHONE: (415) 813-5600
 TELEFAX: (415) 494-0792
 INFORMATION FOR SEQ ID NO: 2:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 5224 base pairs
 TYPE: nucleic acid
 STRANDBNESS: single
 TOPOLOGY: linear
 SEQUENCE DESCRIPTION: SEQ ID NO: 2:
 US-09-898-883-2

Query Match 100.0%; Score 5224; DB 4; Length 5224;
 Best Local Similarity 100.0%; Pred. No: 0; Mismatches 0; Indels 0; Gaps 0;
 Matches 5224; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 GAAATTCTTAGAAATGGGGTAAAGCTGGGTTAGGGGTGGGTGTTAACCCATGGT 60
 1 GAAATTCTTAGAAATGGGGTAAAGCTGGGTTAGGGGTGGGTGTTAACCCATGGT 60
 Db 61 GAGATAGCATGGGTTAATGCTTCACACACATCACTTCATAGAATTAGG 120
 Qy 61 GAGATAGCATGGGTTAATGCTTCACACACATCACTTCATAGAATTAGG 120
 Db 61 GAGATAGCATGGGTTAATGCTTCACACACATCACTTCATAGAATTAGG 120
 Qy 121 AACAGACTATGGCTGGAGGACTGGTGGATCTGGTGTCTGCTGTTACCCCTAGT 180
 Db 121 AACAGACTATGGCTGGAGGACTGGTGGATCTGGTGTCTGCTGTTACCCCTAGT 180
 Qy 181 GTTCTATGGGCTGGTTTAAGCTGGCTCGGGGCTCGGGGTGATATCCTACCTGGTGTCTGCTGTTACCTGGTGTCTGCTGTTACCCCTAGT 180
 Db 241 AGGCCBAGGACTGRCCTCTTCATATCTGTTGGGCTCGGGGTGATATCCTACCTGGTGTCTGCTGTTACCCCTAGT 240
 Qy 241 AGGCCBAGGACTGRCCTCTTCATATCTGTTGGGCTCGGGGTGATATCCTACCTGGTGTCTGCTGTTACCCCTAGT 240
 Db 241 AGGCCBAGGACTGRCCTCTTCATATCTGTTGGGCTCGGGGTGATATCCTACCTGGTGTCTGCTGTTACCCCTAGT 300
 Qy 301 GAGAAATATAACATTCATAAATGTCGGGAGATGAGTGGATGAGATGGTGTCT 360
 Db 301 GAGAAATATAACATTCATAAATGTCGGGAGATGAGTGGATGAGATGGTGTCT 360
 Qy 361 TCATTGTTATCATGATGATGAGGACAAATAGTGTTGAGGATGAGTGGATGAGTGGTGTCT 420
 Db 361 TCATTGTTATCATGATGATGAGGACAAATAGTGTTGAGGATGAGTGGATGAGTGGTGTCT 420
 Qy 421 GAGGGTAGGGTGTGTTCAATTCTTATTTAAGATAACTTTCTTATTTA 480
 Db 421 GAGGGTAGGGTGTGTTCAATTCTTATTTAAGATAACTTTCTTATTTA 480
 Qy 481 TCTCTGACAAATGTTGCGGACCAAGGAATCACAAAGATGAGTGGTGTCT 540
 Db 481 TCTCTGACAAATGTTGCGGACCAAGGAATCACAAAGATGAGTGGTGTCT 540
 Qy 541 AGAAAGTTAAATGTCGAGGAATAATTACATGCTTACAATGACTATGATTTACCAA 600
 Db 541 AGAAAGTTAAATGTCGAGGAATAATTACATGCTTACAATGACTATGATTTACCAA 600
 Qy 601 ACAAGAGTTTCATGAGAAATAATCTGAGGTTAATGGTGTCAAGGGTGGAGGG 660
 Db 601 ACAAGAGTTTCATGAGAAATAATCTGAGGTTAATGGTGTCAAGGGTGGAGGG 660
 Qy 661 CTCTCTCTACTAGAGACTACAGAAATCATCAGGGTAATTATGGTGTCT 720
 Db 661 CTCTCTCTACTAGAGACTACAGAAATCATCAGGGTAATTATGGTGTCT 720
 Qy 721 TAAGGGTTGGGTTACATTTGTCAGGATTGAGGAGGAGGAGGATTTGAAAC 780
 Db 721 TAAGGGTTGGGTTACATTTGTCAGGATTGAGGAGGAGGAGGATTTGAAAC 780
 Qy 781 ATTTTCACTAACCAACCACCAATCCAAACAAACAAAGAAAGAATCTCAGAAC 840
 Db 781 ATTTTCACTAACCAACCACCAATCCAAACAAACAAAGAAAGAATCTCAGAAC 840
 Qy 1801 AGTTACTAGCTTGTGACTTATCTCATCCACCTGCTCATCCACCTATGGGACTT 1860
 Db 1801 AGTTACTAGCTTGTGACTTATCTCATCCACCTGCTCATCCACCTATGGGACTT 1860
 Qy 1861 TTGACAGTATATGGGAAACTCTCTACTGGTGTCTCTATCATGTTATCCCTT 1920
 Db 1861 TTGACAGTATATGGGAAACTCTCTACTGGTGTCTCTATCATGTTATCCCTT 1920
 Qy 1921 TGAACAAAGAGACAGTTCAAAATACAATATGATTATTAGCCCTTGTCT 1980

| | | | | |
|----|------|---|------|----|
| Qy | 1921 | TGAAACAAAGAGAGAAGTTCAATAACATAATTTTATGGCTCCCTTAATCTTTGTG | 1980 | Qy |
| Qy | 1981 | CTATATAGTCGCCAGAGGAGTTAAACTCATTAAAGCTTGAGATGTCGCCCT | 2040 | Qy |
| Db | 1981 | CTATATAGTCGCCAGAGGAGTTAAACTCATTAAAGCTTGAGATGTCGCCCT | 2040 | Db |
| Qy | 2041 | TGCCACTTGCAGGAATTCCAAATCGTAGTTTCTRACTAACTTAACTTGGCCCT | 2100 | Qy |
| Db | 2041 | TGCCACTTGCAGGAATTCCAAATCGTAGTTTCTRACTAACTTAACTTGGCCCT | 2100 | Db |
| Qy | 2101 | TTCAGAACCTCATTCATCGTCACTTCTCTAGTTCTRACTAACTTAACTTGGCCCT | 2160 | Qy |
| Db | 2101 | TTCAGAACCTCATTCATCGTCACTTCTCTAGTTCTRACTAACTTAACTTGGCCCT | 2160 | Db |
| Qy | 2161 | TTTCACCACTTACATTCTCGAACACTAACCCCTCCCTCATTTGCCACC | 2220 | Qy |
| Db | 2161 | TTTCACCACTTACATTCTCGAACACTAACCCCTCCCTCATTTGCCACC | 2220 | Db |
| Qy | 2221 | TCTATTTCAGATCCATAGAGTTAGAGTTACTTCAGGTTACTTCAGGTT | 2280 | Qy |
| Db | 2221 | TCTATTTCAGATCCATAGAGTTAGAGTTACTTCAGGTTACTTCAGGTT | 2280 | Db |
| Qy | 2281 | CCTCCAAAGATGTCAGTCTCCATAGAGTTACTTCAGGTTACTTCAGGTT | 2340 | Qy |
| Db | 2281 | CCTCCAAAGATGTCAGTCTCCATAGAGTTACTTCAGGTTACTTCAGGTT | 2340 | Db |
| Qy | 2341 | TGTTGTTACTGATGACTATTGTTCTCTGATGAGCTGAGCTCAAGAGSC | 2400 | Qy |
| Db | 2341 | TGTTGTTACTGATGACTATTGTTCTCTGATGAGCTGAGCTCAAGAGSC | 2400 | Db |
| Qy | 2401 | TACATGGCTGCTGCTGTTGATGATCCAAAGAGGCTGGATGCA | 2460 | Qy |
| Db | 2401 | TACATGGCTGCTGCTGTTGATGATCCAAAGAGGCTGGATGCA | 2460 | Db |
| Qy | 2461 | GACTAACATTATGTAATTAACCCATCATCGATGTCGACTTGTGCA | 2520 | Qy |
| Db | 2461 | GACTAACATTATGTAATTAACCCATCATCGATGTCGACTTGTGCA | 2520 | Db |
| Qy | 2521 | AGCCCGCTCTGGCATTATTTATGATTATTATCTCATTAACATGAG | 2580 | Qy |
| Db | 2521 | AGCCCGCTCTGGCATTATTTATGATTATTATCTCATTAACATGAG | 2580 | Db |
| Qy | 2581 | GAGGACTACATCACCTATTATGATGAGAAAGCCAGAATGAA | 2640 | Qy |
| Db | 2581 | GAGGACTACATCACCTATTATGATGAGAAAGCCAGAATGAA | 2640 | Db |
| Qy | 2641 | AACCTACCAAAGCTGATGAGTGTGACAGGCCCCAAATTCACCGTTCCA | 2700 | Qy |
| Db | 2641 | AACCTACCAAAGCTGATGAGTGTGACAGGCCCCAAATTCACCGTTCCA | 2700 | Db |
| Qy | 2701 | TTACGTGATTATCTGCTATCTGCTCTGATCATGCGATGGATGAGCAT | 2760 | Qy |
| Db | 2701 | TTACGTGATTATCTGCTATCTGCTCTGATCATGCGATGGATGAGCAT | 2760 | Db |
| Qy | 2761 | CTGCCCGTAAAGGCGAAATGGAGGAGTGGGGATGACACAAACCGCAT | 2820 | Qy |
| Db | 2761 | CTGCCCGTAAAGGCGAAATGGAGGAGTGGGGATGACACAAACCGCAT | 2820 | Db |
| Qy | 2821 | CACAGGAAAGTCAACACAGGACTGAATGAGAAAGTTGTTACTCTG | 2880 | Qy |
| Db | 2821 | CACAGGAAAGTCAACACAGGACTGAATGAGAAAGTTGTTACTCTG | 2880 | Db |
| Qy | 2881 | GCATGACATCTGAGGAAATGGGGCTAACACACAATGGCTATGTCATA | 2940 | Qy |
| Db | 2881 | GCATGACATCTGAGGAAATGGGGCTAACACACAATGGCTATGTCATA | 2940 | Db |
| Qy | 2941 | TTATGTGTTGTTGTTGTTATATATATATGAAATGAAATGCTAT | 3000 | Qy |
| Db | 2941 | TTATGTGTTGTTGTTGTTATATATATGAAATGAAATGCTAT | 3000 | Db |
| Qy | 3001 | TTCTAAAGGCTACATTGCGGCATTTACTTGTCCCTPATCTTGTG | 3060 | Qy |

Qy 4141 GCGCTGTTATACCCCTACTTATGAATGTACCATATTAACTTCTGGTGGGTTA 4200
 Db 4141 GCGCTGTTATACCCCTACTTATGAATGTACCATATTAACTTCTGGTGGGTTA 4200

Qy 4201 TTGATTTTAACTTACCTTAGATAAGAAACCTTGAGACCTTGAACCTTGAATTA 4260
 Db 4201 TTGATTTTAACTTACCTTAGATAAGAAACCTTGAGACCTTGAACCTTGAATTA 4260

Qy 4261 GGTGCCCTGGGTCACCTCCAGATCTGTTAACCTGGGTGAGCTTGAGAATCT 4320
 Db 4261 GGTGCCCTGGGTCACCTCCAGATCTGTTAACCTGGGTGAGCTTGAGAATCT 4320

Qy 4321 TGGGAACTCAAAGTCCCGCAGTGATTCAATGACACCTCTGAGCTTGAATTA 4380
 Db 4321 TGGGAACTCAAAGTCCCGCAGTGATTCAATGACACCTCTGAGCTTGAATTA 4380

Qy 4381 TAGACAGGGATGATAGGGTGGACTCTAGCATCCATTTAAGACAGTGTCAA 440
 Db 4381 TAGACAGGGATGATAGGGTGGACTCTAGCATCCATTTAAGACAGTGTCAA 440

Qy 4441 CATCTGAGTCTAAGTAGTGTGACCTGACCTGACCCACTTATAAAGACAGTGTCAA 4500
 Db 4441 CATCTGAGTCTAAGTAGTGTGACCTGACCTGACCCACTTATAAAGACAGTGTCAA 4500

Qy 4441 CATCTGAGTCTAAGTAGTGTGACCTGACCTGACCCACTTATAAAGACAGTGTCAA 4500

Db 4441 CATCTGAGTCTAAGTAGTGTGACCTGACCTGACCCACTTATAAAGACAGTGTCAA 4500

Qy 4501 TAAATATTATAGGCTTGTGCTCTAAGTGTGTTCTAGTCTGTAGCC 4560
 Db 4501 TAAATATTATAGGCTTGTGCTCTAAGTGTGTTCTAGTCTGTAGCC 4560

Qy 4561 ATACATCGCAACTAACTCTTAAATGTCCTTATGAACTACATTTAGTAAAG 4620
 Db 4561 ATACATCGCAACTAACTCTTAAATGTCCTTATGAACTACATTTAGTAAAG 4620

Qy 4621 TGTTCCTGAGGTAACATGACGACAATTAATCAAGCATATAATTGAAATGATT 4680
 Db 4621 TGTTCCTGAGGTAACATGACGACAATTAATCAAGCATATAATTGAAATGATT 4680

Qy 4681 ATTATCTGCACTAGGACAGTCATCTCTGAAATTGTACTTGAGGATT 4740
 Db 4681 ATTATCTGCACTAGGACAGTCATCTCTGAAATTGTACTTGAGGATT 4740

Qy 4741 GTTATTTGAGAGAGAGGTCTGAACTTCTGTCAGAACAGTCTGTCCTGCA 4800
 Db 4741 GTTATTTGAGAGAGAGGTCTGAACTTCTGTCAGAACAGTCTGTCCTGCA 4800

Qy 4801 TGTAGTCGATTATAGTTTAAAGATAATGATTAGTGTGAACTACCAACT 4860
 Db 4801 TGTAGTCGATTATAGTTTAAAGATAATGATTAGTGTGAACTACCAACT 4860

Qy 4861 GCTGTCTTCTTATGGCTCATACTAAATTGGAGAAATTATTCCTGAACT 4920
 Db 4861 GCTGTCTTCTTATGGCTCATACTAAATTGGAGAAATTATTCCTGAACT 4920

Qy 4921 TAGGGACAAGTCATGTTGAGGAAATTTGTTGAGGAGATTTGTTATGTC 4980
 Db 4921 TAGGGACAAGTCATGTTGAGGAAATTTGTTGAGGAGATTTGTTATGTC 4980

Qy 4981 AAATAATAATAGTTGCAAGGTTTCTGCCAAGAGCTCTGTCCTTGAC 5040
 Db 4981 AAATAATAATAGTTGCAAGGTTTCTGCCAAGAGCTCTGTCCTTGAC 5040

Qy 5041 ATAAATACAAATACCGCTTGCTTAATATGGCAATGTCGATTCAACTA 5100
 Db 5041 ATAAATACAAATACCGCTTGCTTAATATGGCAATGTCGATTCAACTA 5100

Qy 5101 CGAAATACATAAGTACAGATAATCCAAACAAAGGTACTAGTAAACGGCATGCT 5160
 Db 5101 CGAAATACATAAGTACAGATAATCCAAACAAAGGTACTAGTAAACGGCATGCT 5160

Qy 5161 CGAAAGGTAAAGGATTCAGTCAGTTCCATATGGCTCCACACCTGCAAT 5220
 Db 5161 CGAAAGGTAAAGGATTCAGTCAGTTCCATATGGCTCCACACCTGCAAT 5220

Qy 5220 CGAAAGGTAAAGGATTCAGTCAGTTCCATATGGCTCCACACCTGCAAT 5220

Qy 5221 AACAA 5224
 Db 5221 AACAA 5224

Qy US-09-151-376-5
 Db Sequence 5, Application US/09151376

Qy ; Sequence No., 667935

Db ; GENERAL INFORMATION:

Db ; APPLICANT: Henderson, D.R.

Db ; TITLE OF INVENTION: TISSUE SPECIFIC VIRAL VECTORS

Db ; FILE REFERENCE: 34802200021

Db ; CURRENT APPLICATION NUMBER: US/09/151, 376

Db ; CURRENT FILING DATE: 1998-09-10

Db ; EARLIER APPLICATION NUMBER: 08 669, 753

Db ; EARLIER FILING DATE: 1996-06-25

Db ; EARLIER APPLICATION NUMBER: 08 495, 034

Db ; EARLIER FILING DATE: 1995-06-27

Db ; NUMBER OF SEQ ID NOS: 71

Db ; SOFTWARE: Patentin Ver. 2.0

Db ; SEQ ID NO: 5

Db ; LENGTH: 5224

Db ; TYPE: DNA

Db ; ORGANISM: Homo sapiens

Db ; US-09-151-376-5

Db ; Query Match 100.0%; Score 5224; DB 4; Length 5224;

Db ; Best Local Similarity 100.0%; Pred. No: 0; Mismatches 0; Indels 0; Gaps 0;

Db ; Matches 5224; Conservative 0; MiBmatches 0;

Qy 1 GAATCTCTGAAATAGGGGTAGGGTAAATGCTCTCTGTTACCCCATAGT 60
 Db 1 GAATCTCTGAAATAGGGGTAGGGTAAATGCTCTCTGTTACCCCATAGT 60

Qy 61 GAGATAGCATGGGTTAAATGCTTCAACACACATCATTCTAGAAATTAG 120
 Db 61 GAGATAGCATGGGTTAAATGCTTCAACACACATCATTCTAGAAATTAG 120

Qy 121 AACAGACATAGGGCTGGAGACTTGGAGGATGTCCTCTCATACCTGGGTATCT 180
 Db 121 AACAGACATAGGGCTGGAGACTTGGAGGATGTCCTCTCATACCTGGGTATCT 180

Qy 181 GTTCTPATGGGGCTGTTAGCTGCACTTGCAACGGGTCACTGACTCTCCC 240
 Db 181 GTTCTPATGGGGCTGTTAGCTGCACTTGCAACGGGTCACTGACTCTCCC 240

Qy 241 AAGCCCAAGGTACTGCTCTTTCATACTGTTGGGCTCTGGGCTTGATCT 300
 Db 241 AAGCCCAAGGTACTGCTCTTTCATACTGTTGGGCTCTGGGCTTGATCT 300

Qy 301 GAGAAAATAACATTCAATAATGTCCTGTTGAGGATGAGATGAGATGTCAT 360
 Db 301 GAGAAAATAACATTCAATAATGTCCTGTTGAGGATGAGATGAGATGTCAT 360

Qy 361 TCATTGTTGATCATGATGATGAGCAATTAGTGTGAACTTCAACATCT 420
 Db 361 TCATTGTTGATCATGATGATGAGCAATTAGTGTGAACTTCAACATCT 420

Qy 361 TCATTGTTGATCATGATGATGAGCAATTAGTGTGAACTTCAACATCT 420

Db 420 ;

Qy 421 GAGGTAGGGCTGTTACTTCATTAAAGTACTTATCTTCTTATTTA 480
 Db 421 GAGGTAGGGCTGTTACTTCATTAAAGTACTTATCTTCTTATTTA 480

Qy 481 TGCTGTGACAATGTTGTCGGGACCAAGGATCACAGAGTGGCTTGAAATT 540
 Db 481 TGCTGTGACAATGTTGTCGGGACCAAGGATCACAGAGTGGCTTGAAATT 540

Qy 540 ;

Db 540 ;

Qy 541 AAGAAGTTAATGGTCAGGAAATATCATAGTCACAAAGTACTATACCATCA 600
 Db 541 AAGAAGTTAATGGTCAGGAAATATCATAGTCACAAAGTACTATACCATCA 600

Qy 600 ;

Db 600 ;

Qy 601 ACAAGAGGTTCATGAGAAATAATCTGAAGGTTAATAGTGTCAAGGTGAGAGGG 660

| | | | |
|----|------|---|------|
| Db | 601 | ACGAGGGTCATAGAGATAATCTGAGAGTTAATAGTGTCAAGGTGAGGG | 660 |
| Qy | 661 | CTCTCTCTGACTGAGACTAACTCGAAGAACATTAGGTAATATTGAAAGCT | 720 |
| Db | 661 | CTCTCTCTGACTGAGACTAACTCGAAGAACATTAGGTAATATTGAAAGCT | 720 |
| Qy | 721 | TAAGGGTGGGAACTTGTCAGCATGATGAGGAAGGAGGAGTGAATTTGAAAC | 780 |
| Db | 721 | TAAGGGTGGGAACTTGTCAGCATGATGAGGAAGGAGGAGTGAATTTGAAAC | 780 |
| Qy | 781 | ATTTCAGTAACTAACCCACCAACTTCAACAAACAAAGATAAGAAATCTGAAAC | 840 |
| Db | 781 | ATTTCAGTAACTAACCCACCAACTTCAACAAACAAAGATAAGAAATCTGAAAC | 840 |
| Qy | 841 | GTGAGGATAGAGAGGAAATTTCACACACCGTAGTCACTCAGCTCTGAGAA | 900 |
| Db | 841 | GTGAGGATAGAGAGGAAATTTCACACACCGTAGTCACTCAGCTCTGAGAA | 900 |
| Qy | 901 | GTATATATCAATTTAACACTACATCATGCTATAATGATAATTAACCTGATT | 960 |
| Db | 901 | GTATATATCAATTTAACACTACATCATGCTATAATGATAATTAACCTGATT | 960 |
| Qy | 961 | TTPATGCTATAGTACAGGCATTAGAGATTTACCTTATATCAAATA | 1020 |
| Db | 961 | TTPATGCTATAGTACAGGCATTAGAGATTTACCTTATATCAAATA | 1020 |
| Qy | 1021 | ACTTGAGGGATGATCATTTCTGATATGAGAAATTAAACGATGATAT | 1080 |
| Db | 1021 | ACTTGAGGGATGATCATTTCTGATATGAGAAATTAAACGATGATAT | 1080 |
| Qy | 1081 | TTGCTGTATACAGCTAAATGACCATAAGAACATTAGATTTAGTTGAAAT | 1140 |
| Db | 1081 | TTGCTGTATACAGCTAAATGACCATAAGAACATTAGATTTAGTTGAAAT | 1140 |
| Qy | 1141 | CTTCTCTAACTAACAGTCACTGTTACTGTCCTGAGCAATTAGGTTGAACT | 1200 |
| Db | 1141 | CTTCTCTAACTAACAGTCACTGTTACTGTCCTGAGCAATTAGGTTGAACT | 1200 |
| Qy | 1201 | TATGAAAGTAACGGAACTAGATCATGAACTTCTGAAACATGTCGTGAACTCT | 1260 |
| Db | 1201 | TATGAAAGTAACGGAACTAGATCATGAACTTCTGAAACATGTCGTGAACTCT | 1260 |
| Qy | 1261 | ACTTGGACTAACTTGAGCAATACACACATAGATGGGTTGTTGTTGAT | 1320 |
| Db | 1261 | ACTTGGACTAACTTGAGCAATACACACATAGATGGGTTGTTGTTGAT | 1320 |
| Qy | 1321 | ACAACTCTGTCAAAGCTCTTATGCTGTCCTGAAATTGCTGTTCTCAT | 1380 |
| Db | 1321 | ACAACTCTGTCAAAGCTCTTATGCTGTCCTGAAATTGCTGTTCTCAT | 1380 |
| Qy | 1381 | GGTTCTCTTCACTGCTATCTTCTCACCACTCAGTGCTACAATCTGTC | 1440 |
| Db | 1381 | GGTTCTCTTCACTGCTATCTTCTCACCACTCAGTGCTACAATCTGTC | 1440 |
| Qy | 1441 | TGCAAGCTTATGATTCGAAATCTCTCTGAGCTCACTTGTGAGATAAA | 1500 |
| Db | 1441 | TGCAAGCTTATGATTCGAAATCTCTGAGCTCACTTGTGAGATAAA | 1500 |
| Qy | 1501 | AGTAGTATCAATGACATCACTGAGGCTTAAGAGCTTCAACATA | 1560 |
| Db | 1501 | AGTAGTATCAATGACATCACTGAGGCTTAAGAGCTTCAACATA | 1560 |
| Qy | 1561 | CAACCGGGAGTTCTGGATGTTCTAAATGTCCTGAGCATGGGCC | 1620 |
| Db | 1561 | CAACCGGGAGTTCTGGATGTTCTAAATGTCCTGAGCATGGGCC | 1620 |
| Qy | 1621 | TCTGTGCTCTAAACTCTTACTGCCCCACTTGGGAGT | 1680 |
| Db | 1621 | TCTGTGCTCTAAACTCTTACTGCCCCACTTGGGAGT | 1680 |
| Qy | 1681 | AGGTGAAAGGGAGCTGATTAATACAGTCACTGGCATAGGCCAGGACT | 1740 |
| Db | 1681 | AGGTGAAAGGGAGCTGATTAATACAGTCACTGGCATAGGCCAGGACT | 1740 |
| Qy | 1741 | GTTGGGAAACTGACTTCACTTCTAAATATCTCAACATGACATGACTT | 1800 |
| Db | 1741 | GTTGGGAAACTGACTTCACTTCTAAATATCTCAACATGACATGACTT | 1800 |
| Qy | 1801 | AGTACTAAGCTTCTGACTTATCTCATCATTACACTCGCTTATCCGGCCATT | 1860 |
| Db | 1801 | AGTACTAAGCTTCTGACTTATCTCATCATTACACTCGCTTATCCGGCCATT | 1860 |
| Qy | 1861 | TGACGATATTGCGAAACTCTCAACTGCTCATACACTGCTTATCCGGCCATT | 1920 |
| Db | 1861 | TGACGATATTGCGAAACTCTCAACTGCTCATACACTGCTTATCCGGCCATT | 1920 |
| Qy | 1921 | TGAAACAAAGAGCAGTTCAAAATACAAATAGTGTATTAGTGTCTCTGTT | 1980 |
| Db | 1921 | TGAAACAAAGAGCAGTTCAAAATACAAATAGTGTATTAGTGTCTCTGTT | 1980 |
| Qy | 1981 | CTATATAGTCCAGAGGAGTTAAACCTCATTAAGAGTCTTGGAGTGGCCCT | 2040 |
| Db | 1981 | CTATATAGTCCAGAGGAGTTAAACCTCATTAAGAGTCTTGGAGTGGCCCT | 2040 |
| Qy | 2041 | TGCGACTTGTGCGAGAATCCCATATCTGTTCTGTTCTGTTCTGCTC | 2100 |
| Db | 2041 | TGCGACTTGTGCGAGAATCCCATATCTGTTCTGCTC | 2100 |
| Qy | 2101 | TCAAAACTCCATTCTCATCCATTGTCATTGTCCTTACGGTTGTT | 2160 |
| Db | 2101 | TCAAAACTCCATTCTCATCCATTGTCCTTACGGTTGTT | 2160 |
| Qy | 2161 | TTCCACCCCTTACATTTCTGGAAACATTAACCTTACCCCTCTCATTTGGCCACC | 2220 |
| Db | 2161 | TTCCACCCCTTACATTTCTGGAAACATTAACCTTACCCCTCTCATTTGGCCACC | 2220 |
| Qy | 2221 | TCTATTTCTCAGCTCCTGAGATGTTCTCAGGAGTCTTCTCAGGAGCTT | 2280 |
| Db | 2221 | TCTATTTCTCAGCTCCTGAGATGTTCTCAGGAGCTT | 2280 |
| Qy | 2281 | CCTCCAAGATGTCATGACTTCTCTTCTCATCTACTAATCACGATCCACACCA | 2340 |
| Db | 2281 | CCTCCAAGATGTCATGACTTCTCTTCTCATCTACTAATCACGATCCACACCA | 2340 |
| Qy | 2341 | TGTGTGATTACATCATCTATTGTCGTTCTGTTCTGTTGATAGGAGCTAAGTCACAGAGC | 2400 |
| Db | 2341 | TGTGTGATTACATCATCTATTGTCGTTCTGTTGATAGGAGCTAAGTCACAGAGC | 2400 |
| Qy | 2401 | TACATGGTCTGCTGCTGTTCTGTTGATTCATCCAAACAGTCCTGGAATGCA | 2460 |
| Db | 2401 | TACATGGTCTGCTGCTGTTCTGTTGATTCATCCAAACAGTCCTGGAATGCA | 2460 |
| Qy | 2461 | GACTAACATTATGATGAAATAAAACCCATCTCTGAGGCTCTGTGCA | 2520 |
| Db | 2461 | GACTAACATTATGATGAAATAAAACCCATCTCTGAGGCTCTGTGCA | 2520 |
| Qy | 2521 | AGACCGGGTCTGAGGCAATTATGTTGATTAATGCACTATCGCTACTTGTGCA | 2580 |
| Db | 2521 | AGACCGGGTCTGAGGCAATTATGTTGATTAATGCACTATCGCTACTTGTGCA | 2580 |
| Qy | 2581 | GAGCTACTATCTCTTATGTTGATTAATGCTCATGAGGAAATGAA | 2640 |
| Db | 2581 | GAGCTACTATCTCTTATGTTGATTAATGCTCATGAGGAAATGAA | 2640 |
| Qy | 2641 | AACCTACCGAAAGTCATGACTAAGTCAAGGGCAAATTCAACCGTCCCAC | 2700 |
| Db | 2641 | AACCTACCGAAAGTCATGACTAAGTCAAGGGCAAATTCAACCGTCCCAC | 2700 |
| Qy | 2701 | TAGTGTATAATGCTGCTACTGCTCTGATCATGAGTCAGACAT | 2760 |
| Db | 2701 | TAGTGTATAATGCTGCTACTGCTCTGATCATGAGTCAGACAT | 2760 |
| Qy | 2761 | CTCTCTCTAAAGCAGAATATGAGGAGTTGAGGAGTCAACACAGCAATAT | 2820 |
| Db | 2761 | CTCTCTCTAAAGCAGAATATGAGGAGTTGAGGAGTCAACACAGCAATAT | 2820 |

| | | | | | | | |
|----|------|--|------|----|------|--|------|
| Qy | 2821 | CAGAGGAAAGTCCAAAACAGSACCTGACTGTAGAAAGTGTTACTCCGGGTAGTC | 2880 | Qy | 3901 | AGATGATAATTCACTTACATCTAGAGAACCAAATGTTAAACTTGTTG | 3960 |
| Db | 2821 | CAGAGGAAACTCCAAACAGSACCTGACTGTAGAAAGTGTTACTCCGGGTAGTC | 2880 | Db | 3901 | AGATGATAATTCACTTACATCTAGAGAACCAAATGTTAAACTTGTTG | 3960 |
| Qy | 2881 | GCATGCACATCTGTGAACTGGTGTGACAGACACATCTGGTTGTTGTCATA | 2940 | Qy | 3961 | TCCATTATTCCTGTTTCACTCATTTTTAGGGGGGGAGGAAATCAGAGGAG | 4020 |
| Db | 2881 | GCATGCACATCTGTGAACTGGTGTGACAGACACATCTGGTTGTTGTCATA | 2940 | Db | 3961 | TCCATTATTCCTGTTTCACTCATTTTTAGGGGGGGAGGAAATCAGAGGAG | 4020 |
| Qy | 2941 | TATTTGTGTTGTTGTTGTTGTTGTTGTTGTTGTTGTTGTTGTTGTTGTTGTTG | 3000 | Qy | 4021 | GTACATGATAACAAATGAGGACTCTCCATGTTGTTGTTGTTGTTGTTGTTG | 4080 |
| Db | 2941 | TATTTGTGTTGTTGTTGTTGTTGTTGTTGTTGTTGTTGTTGTTGTTGTTGTTG | 3000 | Db | 4021 | GTACATGATAACAAATGAGGACTCTCCATGTTGTTGTTGTTGTTGTTGTTG | 4080 |
| Qy | 3061 | AGATGATCAGTTGATTACTGGCTTGAGATGATATATCTACATCTATATCTAC | 3120 | Qy | 4081 | ACAATATTATGAGCATTTCCATTCTAACATCCACAGTTGTTGAT | 4140 |
| Db | 3061 | AGATGATCAGTTGATTACTGGCTTGAGATGATATATCTACATCTATATCTAC | 3120 | Db | 4081 | ACAATATTATGAGCATTTCCATTCTAACATCCACAGTTGTTGAT | 4140 |
| Qy | 3121 | TATATCTATCTATCTATCTATCTATCTATCTATCTATCTATCTATCTATCTA | 3180 | Qy | 4141 | GGCTTATACACCTTATGATGTTGACACTCTCCATGTTGAAAGCTG | 4200 |
| Db | 3121 | TATATCTATCTATCTATCTATCTATCTATCTATCTATCTATCTATCTATCTA | 3180 | Db | 4141 | GGCTTATACACCTTATGATGTTGACACTCTCCATGTTGAAAGCTG | 4200 |
| Qy | 3241 | GTAACCCCTTCTCTACATTAACCCATGTTGAAACCAATCATTTTCTACATC | 3300 | Qy | 4261 | GGTGCCTGGCTCTCAACTCCACAGTTGTTGAAACCCCTGGAATTG | 4320 |
| Db | 3241 | GTAACCCCTTCTCTACATTAACCCATGTTGAAACCAATCATTTTCTACATC | 3300 | Db | 4261 | GGTGCCTGGCTCTCAACTCCACAGTTGTTGAAACCCCTGGAATTG | 4320 |
| Qy | 3301 | AATGTGCTGAGTCTGGCTATGAACTGAGCAGACTGTAAGAACGCTTGGGATTTGCC | 3360 | Qy | 4321 | TTGGGATTCAAAAGTCCCGACTGATCTAATGTTGAGCTTGTGAAAGCTG | 4380 |
| Db | 3301 | AATGTGCTGAGTCTGGCTATGAACTGAGCAGACTGTAAGAACGCTTGGGATTTGCC | 3360 | Db | 4321 | TTGGGATTCAAAAGTCCCGACTGATCTAATGTTGAGCTTGTGAAAGCTG | 4380 |
| Qy | 3361 | CATGCTTGGGAAGCTTATAGTTGCTTAAACTCTATTCTAGTTCTCAACT | 3420 | Qy | 4381 | TAGACGGGATGAGGTTGAGCTTGTGAACTGCTTGTGAAAGCTG | 4440 |
| Db | 3361 | CATGCTTGGGAAGCTTATAGTTGCTTAAACTCTATTCTAGTTCTCAACT | 3420 | Db | 4381 | TAGACGGGATGAGGTTGAGCTTGTGAACTGCTTGTGAAAGCTG | 4440 |
| Qy | 3421 | AATACTCTACATCTATGTTCTAGTTCTCTACATCTATCTATCTATCTA | 3480 | Qy | 4441 | CATCTGAGTGTCTAATGATGTTGCTTAACTCTAGTTCTCAACT | 4500 |
| Db | 3421 | AATACTCTACATCTATGTTCTAGTTCTCTACATCTATCTATCTATCTA | 3480 | Db | 4441 | CATCTGAGTGTCTAATGATGTTGCTTAACTCTAGTTCTCAACT | 4500 |
| Qy | 3481 | TATTTGAGTAAGTCCCCTTGGAGAGACTGACCTTGTAACTATCC | 3540 | Qy | 4501 | TAATAATTATAGACCTTGTGTTTATAGGCTTGTGCTCTAAGTTCTATGTTAGCC | 4560 |
| Db | 3481 | TATTTGAGTAAGTCCCCTTGGAGAGACTGACCTTGTAACTATCC | 3540 | Db | 4501 | TAATAATTATAGACCTTGTGTTTATAGGCTTGTGCTCTAAGTTCTATGTTAGCC | 4560 |
| Qy | 3540 | TATTTGAGTAAGTCCCCTTGGAGAGACTGACCTTGTAACTATCC | 3540 | Qy | 4561 | ATACATGCTACATCTAACTCTAACTTAAAGTACCTTGTGAAAG | 4620 |
| Db | 3540 | TATTTGAGTAAGTCCCCTTGGAGAGACTGACCTTGTAACTATCC | 3540 | Db | 4561 | ATACATGCTACATCTAACTCTAAAGTACCTTGTGAAAG | 4620 |
| Qy | 3541 | TGGAATCCAAAAGTAGACAGGATGGTGTACCTTGTGAGAGTGGAGCAAG | 3600 | Qy | 4621 | TGTTCTGACCTAACATCACACATTAATCTACAGCATGTTGAAAGTAA | 4680 |
| Db | 3541 | TGGAATCCAAAAGTAGACAGGATGGTGTACCTTGTGAGAGTGGAGCAAG | 3600 | Db | 4621 | TGTTCTGACCTAACATCACACATTAATCTACAGCATGTTGAAAGTAA | 4680 |
| Qy | 3601 | GCCTGTTGTTGTTACACTGTTCTTAGGAGAACCTTAACTGAGAACCTAGCAA | 3660 | Qy | 4681 | ATTATCTGCACTTAGGGAACGTCTCTCTGAAATTGTTGACTTTGAGAGTT | 4740 |
| Db | 3601 | GCCTGTTGTTGTTACACTGTTCTTAGGAGAACCTTAACTGAGAACCTAGCAA | 3660 | Db | 4681 | ATTATCTGCACTTAGGGAACGTCTCTCTGAAATTGTTGACTTTGAGAGTT | 4740 |
| Qy | 3661 | ATGCGAACAAACTACAACTACAACTACAACTACAACTACAACTACAACTACAA | 3720 | Qy | 4741 | GTATATTGCAAGTAGAGGAGCTGATGTTGTCAGTGTGTCAGTGTGTCGCTGCA | 4800 |
| Db | 3661 | ATGCGAACAAACTACAACTACAACTACAACTACAACTACAACTACAACTACAA | 3720 | Db | 4741 | GTATATTGCAAGTAGAGGAGCTGATGTTGTCAGTGTGTCAGTGTGTCGCTGCA | 4800 |
| Qy | 3721 | TATACATGATTAATGTTGTTGTTGAACTGAGCTTGTGAACTTGTGAACT | 3780 | Qy | 4801 | TGATAGGCAATTAACTGTTGAACTGAGCTTGTGAACTTGTGAACT | 4860 |
| Db | 3721 | TATACATGATTAATGTTGTTGTTGAACTGAGCTTGTGAACTTGTGAACT | 3780 | Db | 4801 | TGATAGGCAATTAACTGTTGAACTGAGCTTGTGAACTTGTGAACT | 4860 |
| Qy | 3781 | TAATCTGTCCTGCTAGTAGCTGCACTGATGAACTTTTAACTGTTGACTCTAGCTTA | 3840 | Qy | 4861 | GCTGTTCTTCTTATGCTCTATTAATGTTGAGAGAAATTAACTTCTCACT | 4920 |
| Db | 3781 | TAATCTGTCCTGCTAGTAGCTGCACTGATGAACTTTTAACTGTTGACTCTAGCTTA | 3840 | Db | 4861 | GCTGTTCTTCTTATGCTCTATTAATGTTGAGAGAAATTAACTTCTCACT | 4920 |
| Qy | 3841 | TAAATGTTGCTCTAGTAGCTGCACTGATGTTGTTGAGAGTTGAGAGAG | 3900 | Qy | 4921 | TAGGGCAAGTCATGCTTGTGAACTTCTGAGAGAAATTAACTTCTCACT | 4980 |
| Db | 3841 | TAAATGTTGCTCTAGTAGCTGCACTGATGTTGTTGAGAGTTGAGAGAG | 3900 | Db | 4921 | TAGGGCAAGTCATGCTTGTGAACTTCTGAGAGAAATTAACTTCTCACT | 4980 |
| Qy | 4981 | AAAATAAAATAGTTGCAAGTTTCTGCCAACAGAGCTCTGGTCCCTGAAAC | 5040 | Qy | 4981 | AAAATAAAATAGTTGCAAGTTTCTGCCAACAGAGCTCTGGTCCCTGAAAC | 5040 |

RESULT 6
US-09-151-376-45
; Sequence 45, Application US/09151376
; GENERAL INFORMATION:
; Patent No. 6176935
; APPLICANT: Henderson, D.R.
; TITLE OF INVENTION: TISSUE SPECIFIC VIRAL VECTORS
FILE REFERENCE: 348022000221
CURRENT APPLICATION NUMBER: US/09/151,376
CURRENT FILING DATE: 1998-05-10
; EARLIER APPLICATION NUMBER: 08/169,753
; EARLIER FILING DATE: 1990-06-26
; EARLIER APPLICATION NUMBER: 08/495,034
; EARLIER FILING DATE: 1995-06-27
NUMBER OF SEQ ID NOS: 71
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO: 45
LENGTH: 5224
TYPE: DNA
ORGANISM: Unknown
FEATURE:
; OTHER INFORMATION: Description of Unknown Organism: unknown
; US-09-151-376-45

Query Match 100.0% Score 5224; DB 4; Length 5224;
Best Local Similarity 100.0%; Pred. No. 0; Mismatches 0; Indels 0; Gaps 0;
Matches 5224; Conservative 0;

Qy 1 GRATTCCTTAGAATATGGGGTAGGGGGTGGGGTAATTCTGTGTTACCCCATAGT 60
Db 1 GATTCTCTAGAAATATGGGGTAGGGGGTGGGGTAATTCTGTGTTACCCCATAGT 60

Qy 61 GAGATAGCGATGGTTAACATGCTTCACACACATCACATTCTAGAATAGG 120
Db 61 GGATAGCGATGGTTAACATGCTTCACACACATCACATTCTAGAATAGG 120

Qy 121 AACAGACTATGGCTGGGCTGAGGCTTGAGGATCTGTCATAACACTTGTGATCT 180
Db 121 AACAGACTATGGCTGGGCTGAGGCTTGAGGATCTGTCATAACACTTGTGATCT 180

Qy 181 GCTCTATGGGGCTGGTTAGCTTGCACAGGGTCACTGTTCTGCC 240
Db 181 GCTCTATGGGGCTGGTTAGCTTGCACAGGGTCACTGTTCTGCC 240

Qy 241 AAGCCAAAGGACTGCTCTTCATATCTGCTGGGCTCTGGGCTGAAATCT 300
Db 241 AAGCCAAAGGACTGCTCTTCATATCTGCTGGGCTCTGGGCTGAAATCT 300

Qy 301 GAGAAATATAACTTCAATATCTGCTGGGCTCTGGGCTGAAATCT 360
Db 301 GAGAAATATAACTTCAATATCTGCTGGGCTCTGGGCTGAAATCT 360

Qy 361 TCATTTCTCATGAACTGAGGCAATTAGCTGATAATCCTTAGTACACATCT 420
Db 361 TCATTTCTCATGAACTGAGGCAATTAGCTGATAATCCTTAGTACACATCT 420

Qy 421 GAGGGTGGGGTGGTCAATTCTATTAAAGTACTTCTATTATTAA 480
Db 421 GAGGGTGGGGTGGTCAATTCTATTAAAGTACTTCTATTATTAA 480

Qy 481 TGCTGGGACAATGTTGTOGGGACACAGGAATCAAAGATGTCITGAACT 540
Db 481 TGCTGGGACAATGTTGTOGGGACACAGGAATCAAAGATGTCITGAACT 540

Qy 541 AAGAACTTAATGTCAGGAAATTCATAGCTTACAGTATACCTCAA 600
Db 541 AAGAACTTAATGTCAGGAAATTCATAGCTTACAGTATACCTCAA 600

Qy 601 ACAGAGGCTCATGAGAAATAATCTGAAGGTTATAGTTGTCAGAGGG 660
Db 601 ACAGAGGCTCATGAGAAATAATCTGAAGGTTATAGTTGTCAGAGGG 660

Qy 661 CTCCTCTAGTAGAGACTATCAGAAATACCTGGGATAATTCTGATAGCCT 720
Db 661 CTCCTCTAGTAGAGACTATCAGAAATACCTGGGATAATTCTGATAGCCT 720

Qy 721 TRAGGGTGGGTAATTGTCACAGGTTCAAGGATTGAGGAGGAGGTAATTGAAAC 780
Db 721 TRAGGGTGGGTAATTGTCACAGGTTCAAGGATTGAGGAGGAGGTAATTGAAAC 780

Qy 781 ATTTCACACTACCAACACCACAAACAAACAAAATGAAAGAACTCAGAAC 840
Db 781 ATTTCACACTACCAACACCACAAACAAACAAAATGAAAGAACTCAGAAC 840

Qy 841 GTGAGATAGAGAGGATTCTCAACCCACAGGATAGCTCACTGCTCTGAGAA 900
Db 841 GTGAGATAGAGAGGATTCTCAACCCACAGGATAGCTCACTGCTCTGAGAA 900

Qy 901 GTATATCTTAATTAACTAAGTACGATCATGCTTAATATGATAATATRACTGCTT 960
Db 901 GTATATCTTAATTAACTAAGTACGATCATGCTTAATATGATAATATRACTGCTT 960

Qy 961 TTAACTCTAATGACCGGCTTAGAGATAATTCCATTATATATCMAATAA 1020
Db 961 TTAACTCTAATGACCGGCTTAGAGATAATTCCATTATATCMAATAA 1020

Qy 1021 ACTTGGGGTAGATCTATTCTACATGAAATTGAGAAATTAAACAGATGATT 1080
Db 1021 ACTTGGGGTAGATCTATTCTACATGAAATTGAGAAATTAAACAGATGATT 1080

Qy 1081 TTGCTCTCATACAGCTTAATTGACCATAGAAATTAAACAGATGATT 1140
Db 1081 TTGCTCTCATACAGCTTAATTGACCATAGAAATTAAACAGATGATT 1140

Qy 1141 CTTCATACACAGCTGTTACTGTCATGCTCTGAGCTCAAGACT 1200
Db 1141 CTTCATACACAGCTGTTACTGTCATGCTCTGAGCTCAAGACT 1200

Qy 1201 TATGAAAGTAACGAACTACATCAATGAAAGCATGTCATGAACT 1260
Db 1201 TATGAAAGTAACGAACTACATCAATGAAAGCATGTCATGAACT 1260

Qy 1260 1261 ACTTAGGACTAACATTGAGGATCACTGAACTGAGGATGTTCTGTA 1320
Db 1261 ACTTAGGACTAACATTGAGGATCACTGAACTGAGGATGTTCTGTA 1320

Qy 1320 1321 ACAACTCTGCTCAAGCTCTCTTATGCTCTGAGAAATTGCTGTTCTCAT 1380
Db 1321 ACAACTCTGCTCAAGCTCTCTTATGCTCTGAGAAATTGCTGTTCTCAT 1380

Qy 1380 1381 GGTTCCTTCACTGCTATCTATTTCTCACCTCACATGCTCAACTGTC 1440
Db 1381 GGTTCCTTCACTGCTATCTATTTCTCACCTCACATGCTCAACTGTC 1440

Qy 1440 1441 TCGAACCTTATGATTCCAAATATCTCTAGCTCACTGTTCCAGAGATAAA 1500

| | | | | |
|----|------|---|------|----|
| Db | 1441 | TGCAAGCTATGAACTCCCAAATTTCTACGCTCACTTGTCAGAGATAAAA | 1500 | Qy |
| Qy | 1501 | AGTGTATCAATGCACATCACGCCACTTGAGGGCTTAAGAGCTTCACATA | 1560 | Db |
| Db | 1501 | AGTGTATCAATGCACATCACGCCACTTGAGGGCTTAAGAGCTTCACATA | 1560 | Qy |
| Qy | 1561 | CAAAACGGGGAGTTTGCTGGGAGTTTCCTAAAGAATGTCCTGAGGGTC | 1620 | Db |
| Db | 1561 | CAAAACGGGGAGTTTGCTGGGAGTTTCCTAAAGAATGTCCTGAGGGTC | 1620 | Qy |
| Qy | 1621 | TCTGTGCTCTTAATATCTATTACTTGTGCGCTGTCAGCTGAGGAGATG | 1680 | Db |
| Db | 1621 | TCTGTGCTCTTAATATCTATTACTTGTGCGCTGTCAGCTGAGGAGATG | 1680 | Qy |
| Qy | 1681 | AGAGTGAAGGGAGGCTGTTAAATTAATACACTAGTCATGGCATAGCCGG | 1740 | Db |
| Db | 1681 | AGAGTGAAGGGAGGCTGTTAAATTAATACACTAGTCATGGCATAGCCGG | 1740 | Qy |
| Qy | 1741 | GTTGGGTAACCTGTCACTTTAACTTAACTAATTCAGTCATGGCATAGCCGG | 1800 | Db |
| Db | 1741 | GTTGGGTAACCTGTCACTTTAACTTAACTAATTCAGTCATGGCATAGCCGG | 1800 | Qy |
| Qy | 1801 | AGTGTACTAGTCCTTGACTTATCTCATTCAACCTCAGTTATCCAGGCCTT | 1860 | Db |
| Db | 1801 | AGTGTACTAGTCCTTGACTTATCTCATTCAACCTCAGTTATCCAGGCCTT | 1860 | Qy |
| Qy | 1861 | TGAGAAGTATATGGAAACTCTCAACTGGTCTCTATCATAGTCCTATCCC | 1920 | Db |
| Db | 1861 | TGAGAAGTATATGGAAACTCTCAACTGGTCTCTATCATAGTCCTATCCC | 1920 | Qy |
| Qy | 1921 | TGAAACAAAGAGGAGGTTCAAAATACAAATAGTTATAGCTCCCTTTGTT | 1980 | Db |
| Db | 1921 | TGAAACAAAGAGGAGGTTCAAAATACAAATAGTTATAGCTCCCTTTGTT | 1980 | Qy |
| Qy | 1981 | CTTAAATAGTCCTCAAGAGGTTATAACTCCATAGTCCTTATGCTCCCTT | 2040 | Db |
| Db | 1981 | CTTAAATAGTCCTCAAGAGGTTATAACTCCATAGTCCTTATGCTCCCTT | 2040 | Qy |
| Qy | 2041 | TGCAACTTGCCAGGAATCCCATAATCTAGTTCTACTTAACTTGCCCTC | 2100 | Db |
| Db | 2041 | TGCAACTTGCCAGGAATCCCATAATCTAGTTCTACTTAACTTGCCCTC | 2100 | Qy |
| Qy | 2101 | TTCAAACTGCAATTCTCTCATCCCTTAAGTGCGATGTTCCCTAACCGGTTG | 2160 | Db |
| Db | 2101 | TTCAAACTGCAATTCTCTCATCCCTTAAGTGCGATGTTCCCTAACCGGTTG | 2160 | Qy |
| Qy | 2161 | TTTCAACCACCTTTACATTCTCCATTCCTTAAGTGCGATGTTCCCTAACCGGTTG | 2220 | Db |
| Db | 2161 | TTTCAACCACCTTTACATTCTCCATTCCTTAAGTGCGATGTTCCCTAACCGGTTG | 2220 | Qy |
| Qy | 2221 | TCTAAATTCTTCGATCTCCATGAGGAGTGTACTTCCTCCAGGAAGGCCATTCTGAC | 2280 | Db |
| Db | 2221 | TCTAAATTCTTCGATCTCCATGAGGAGTGTACTTCCTCCAGGAAGGCCATTCTGAC | 2280 | Qy |
| Qy | 2281 | CCTCCAAAGATGTCATGTTTACATTCTACTATACCTCCCTTCATTCGCCACC | 2340 | Db |
| Db | 2281 | CCTCCAAAGATGTCATGTTTACATTCTACTATACCTCCCTTCATTCGCCACC | 2340 | Qy |
| Qy | 2341 | TGTTGTGATTACTGATCATGTTCTCTTGATGAGCTAACCTCAACAGAGC | 2400 | Db |
| Db | 2341 | TGTTGTGATTACTGATCATGTTCTCTTGATGAGCTAACCTCAACAGAGC | 2400 | Qy |
| Qy | 2401 | TACATGTTCTGTCCTGTTCTGTTCTGATGAGCTAACCTCAACAGAGC | 2460 | Db |
| Db | 2401 | TACATGTTCTGTCCTGTTCTGATGAGCTAACCTCAACAGAGC | 2460 | Qy |
| Qy | 2461 | GACTTAACTTATGATGAAATAACCCCATCTATGAGTCTGACTTGTC | 2520 | Db |
| Db | 2461 | GACTTAACTTATGATGAAATAACCCCATCTATGAGTCTGACTTGTC | 2520 | Qy |
| Qy | 2521 | AGACCGGTCTGAGGCAATTAAATTGATTTAATCTCATTAACCATGAG | 2580 | Db |
| Db | 2521 | AGACCGGTCTGAGGCAATTAAATTGATTTAATCTCATTAACCATGAG | 2580 | Qy |
| Qy | 2581 | GAGGTTACTACATCTCTTATAGTGTAAAGATAAGCCAGAGTAATT | 2640 | Db |
| Db | 2581 | GAGGTTACTACATCTCTTATAGTGTAAAGATAAGCCAGAGTAATT | 2640 | Qy |
| Qy | 2641 | ACTTACCAAAGTCATGTTAGTGTAAAGGGCAAATTCAACAGTCCTCA | 2700 | Db |
| Db | 2641 | ACTTACCAAAGTCATGTTAGTGTAAAGGGCAAATTCAACAGTCCTCA | 2700 | Qy |
| Qy | 2701 | TACGTGTTATACTGTGCTATCTGCCTCTGATCATATGGCATGATGAGACAT | 2760 | Db |
| Db | 2701 | TACGTGTTATACTGTGCTATCTGCCTCTGATCATATGGCATGATGAGACAT | 2760 | Qy |
| Qy | 2761 | CTGCRCCGTTAGGAGAAATGGAGGAGATGGAGGATGAGACAT | 2820 | Db |
| Db | 2761 | CTGCRCCGTTAGGAGAAATGGAGGATGAGACAT | 2820 | Qy |
| Qy | 2821 | CAGAGAAAGCCAAACAGGACTGAACTGAGAAGTGTGACTCTGGTGACT | 2880 | Db |
| Db | 2821 | CAGAGAAAGCCAAACAGGACTGAACTGAGAAGTGTGACTCTGGTGACT | 2880 | Qy |
| Qy | 2941 | TATTTGTGTTGTTGTTATATATATATATATATATATATATATATATAT | 3000 | Db |
| Db | 2941 | TATTTGTGTTGTTGTTATATATATATATATATATATATATATATATAT | 3000 | Qy |
| Qy | 3001 | TACTAAAGCCCTACATTGGCCAGGATTTCATGTTGCCCCCTAACTTTGAG | 3060 | Db |
| Db | 3001 | TACTAAAGCCCTACATTGGCCAGGATTTCATGTTGCCCCCTAACTTTGAG | 3060 | Qy |
| Qy | 3061 | AGATGTCAGATGGATTACTGGCCCTTGAGATGATATCTACATCTATCTAC | 3120 | Db |
| Db | 3061 | AGATGTCAGATGGATTACTGGCCCTTGAGATGATATCTACATCTATCTAC | 3120 | Qy |
| Qy | 3121 | TATATCTATCTATCTATCTATCTATCTATCTATCTATCTATCTATCTATCT | 3180 | Db |
| Db | 3121 | TATATCTATCTATCTATCTATCTATCTATCTATCTATCTATCTATCT | 3180 | Qy |
| Qy | 3181 | AATATGTTGTAAGTTAAAGATTGAGCTTATAGATCTGGATTTCGAAT | 3240 | Db |
| Db | 3181 | AATATGTTGTAAGTTAAAGATTGAGCTTATAGATCTGGATTTCGAAT | 3240 | Qy |
| Qy | 3241 | GTAACCCCTTCTCTACATTAACCCATGTTGAGCTTATAGATCTGGATTTCGAAT | 3300 | Db |
| Db | 3241 | GTAACCCCTTCTCTACATTAACCCATGTTGAGCTTATAGATCTGGATTTCGAAT | 3300 | Qy |
| Qy | 3301 | AAATGTGCTGAGCTCTGCTGCTGATGAGAGCACTGTGAAAGCCATTGGATTTGCC | 3360 | Db |
| Db | 3301 | AAATGTGCTGCTGCTGCTGATGAGAGCACTGTGAAAGCCATTGGATTTGCC | 3360 | Qy |
| Qy | 3361 | CATGTTGGCCGAAGCTTATAGTTGCTCTTCAAAACTCTTTCAGTTGCTCTAATC | 3420 | Db |
| Db | 3361 | CATGTTGGCCGAAGCTTATAGTTGCTCTTCAAAACTCTTTCAGTTGCTCTAATC | 3420 | Qy |
| Qy | 3421 | AATACTCTAGCTATGCTTCTGTTCTGATGAGCTAACCTTTCATACATTGCTCTAATC | 3480 | Db |
| Db | 3421 | AATACTCTAGCTATGCTTCTGATGAGCTAACCTTTCATACATTGCTCTAATC | 3480 | Qy |
| Qy | 3481 | TATTTGAGTAAGTCCCTTGAGGAAGTAGAGACTGCACTTGTAAACTATCC | 3540 | Db |
| Db | 3481 | TATTTGAGTAAGTCCCTTGAGGAAGTAGAGACTGCACTTGTAAACTATCC | 3540 | Qy |
| Qy | 3541 | TGGATCCAAAGGATAGACAAGGATGGCTACTCTCTGAGGAGTACGTGAGCAG | 3600 | Db |
| Db | 3541 | TGGATCCAAAGGATAGACAAGGATGGCTACTCTCTGAGGAGTACGTGAGCAG | 3600 | Qy |
| Qy | 3601 | GGCTTTGTTGTTAGATGTTCTGAGGAGCAACTAGGAGAGCAGCATGAG | 3660 | Db |
| Db | 3601 | GGCTTTGTTGTTAGATGTTCTGAGGAGCAACTAGGAGAGCAGCATGAG | 3660 | Qy |

| | | | | |
|---|------|--|------|----|
| Y | 121 | AACAGACTATGGCTGAGGACTTGTGAGATGTCGTCTCATACACTGGGTGATCT | 180 | Qy |
| b | 121 | AACAGACTATGGCTGAGGACTTGTGAGATGTCGTCTCATACACTGGGTGATCT | 180 | Db |
| b | 181 | GTTCTATGGGCTGTTAACGCTTGCGCAACTTGCAACAGGGTCACTGACTTCTCCC | 240 | Qy |
| b | 241 | AAGCCAGGACTACTGCTCTTCAATCTGTTGGGCTCTGGGCTGAATCT | 300 | Db |
| b | 241 | AAGCCAGGACTACTGCTCTTCAATCTGTTGGGCTCTGGGCTGAATCT | 300 | Qy |
| b | 241 | AAGCCAGGACTACTGCTCTTCAATCTGTTGGGCTCTGGGCTGAATCT | 300 | Db |
| b | 301 | GAGAAATTAAACATCAATGTCGTGAGTAGTGTGAGTGTGCT | 360 | Qy |
| b | 301 | GAGAAATTAAACATCAATGTCGTGAGTAGTGTGAGTGTGCT | 360 | Db |
| b | 361 | TGATTGTTATCATGTAATGAAATGGGCAATTAGTGATAAATCCTTGTTAGTACACACT | 420 | Qy |
| b | 421 | GAGGGTAGGGGTGACTATTCAATTCTATTATAAGATACTTCTATTATTA | 480 | Db |
| b | 421 | GAGGGTAGGGGTGACTATTCAATTCTATTATAAGATACTTCTATTATTA | 480 | Qy |
| b | 481 | TGCTTGACAAATGTTGTGGACACAGGATCAAAGATGAGCTTGTGATT | 540 | Db |
| b | 481 | TGCTTGACAAATGTTGTGGACACAGGATCAAAGATGAGCTTGTGATT | 540 | Qy |
| b | 541 | AAGAAAGTTAATGGTCCGGAAATAATTACATAGTTACAAATGACTATATACCA | 600 | Db |
| b | 601 | ACAAGGGTCCATGACAAATAATCTGAAGGTTAAAGGTAAAGGAGG | 600 | Qy |
| b | 601 | ACAAGGGTCCATGACAAATAATCTGAAGGTTAAAGGTAAAGGAGG | 600 | Db |
| b | 61 | CTCTCTCTAGTAGAGACTAACGAAATCATCGGGATAATTGAAATAGCT | 720 | Qy |
| b | 61 | CTCTCTCTAGTAGAGACTAACGAAATCATCGGGATAATTGAAATAGCT | 720 | Db |
| b | 721 | TAAGGGTGGTACATTGTCAGGATGAGGAGGAGGAGGATATGAAAC | 780 | Qy |
| b | 721 | TAAGGGTGGTACATTGTCAGGATGAGGAGGAGGAGGATATGAAAC | 780 | Db |
| b | 781 | ATTTCAACTACCAACCAACATCCACAAACAAACAAATGAAAGAACTCGAAC | 840 | Qy |
| b | 781 | ATTTCAACTACCAACCAACATCCACAAACAAATGAAAGAACTCGAAC | 840 | Db |
| b | 841 | GTGACATAGAGAAGGATTCACGACACCTATGCTCACTGCTGAAAGA | 900 | Qy |
| b | 841 | GTGACATAGAGAAGGATTCACGACACCTATGCTCACTGCTGAAAGA | 900 | Db |
| b | 901 | GTATACATATTAACACTACATCTGTAATATGATAATATATGCTT | 960 | Qy |
| b | 901 | GTATACATATTAACACTACATCTGTAATATGATAATATATGCTT | 960 | Db |
| b | 961 | TTAATGCTATAAGTACAGGCAATTAGAACATCTGTAATATGATAATATGCTT | 1020 | Qy |
| b | 961 | TTAATGCTATAAGTACAGGCAATTAGAACATCTGTAATATGATAATATGCTT | 1020 | Db |
| b | 1021 | ACTGAGGGTAGATCTTCATGATATGAGAAAATTAACAGATGAAAT | 1080 | Qy |
| b | 1021 | ACTGAGGGTAGATCTTCATGATATGAGAAAATTAACAGATGAAAT | 1080 | Db |
| b | 1081 | TGCTCTCATACAGCTATAATGACGATAGAACATAGATTAATGTTGATT | 1140 | Qy |
| b | 1081 | TGCTCTCATACAGCTATAATGACGATAGAACATAGATTAATGTTGATT | 1140 | Db |
| b | 1141 | CTTCTCATACAGCTATAATGACGATAGAACATAGATTAATGTTGATT | 1200 | Qy |
| b | 1141 | CTTCTCATACAGCTATAATGACGATAGAACATAGATTAATGTTGATT | 1200 | Db |
| b | 1201 | TATGAAAGTAACCGATCGAAATCACATCGAAACCATCTGAGACTCTG | 1260 | Qy |
| b | 1201 | TATGAAAGTAACCGATCGAAATCACATCGAAACCATCTGAGACTCTG | 1260 | Db |
| b | 1261 | ACTTAGACTAACTTGAGGATACACAGATGTTGAGGATGTTGCTTAGCT | 1320 | Qy |
| b | 1321 | ACAACTCTGTTCAAGCTCTCTTATGCTGCTTGAATTTGCTTCTCAT | 1380 | Db |
| b | 1321 | ACAACTCTGTTCAAGCTCTCTTATGCTGCTTGAATTTGCTTCTCAT | 1380 | Qy |
| b | 1381 | GGTTCTCTTCACTGCTATCTTCTCAACCTCACATGCCTACATACAGTC | 1440 | Db |
| b | 1381 | GGTTCTCTTCACTGCTATCTTCTCAACCTCACATGCCTACATACAGTC | 1440 | Qy |
| b | 1441 | TGCAAGCTTATGATCCAAATATCTAATCTGCTAGCCCAATTGTTGCTCAGAGATAAA | 1500 | Db |
| b | 1441 | TGCAAGCTTATGATCCAAATATCTAATCTGCTAGCCCAATTGTTGCTCAGAGATAAA | 1500 | Qy |
| b | 1501 | AGTAGATTCATGATCCAAATATCTAATCTGCTAGCCCAATTGTTGCTCAGAGATAAA | 1560 | Db |
| b | 1501 | AGTAGATTCATGATCCAAATATCTAATCTGCTAGCCCAATTGTTGCTCAGAGATAAA | 1560 | Qy |
| b | 1561 | CAACGGGGAGGTTGCTGAAAGTTCTCAAATGTCGCTGTAGCACTAGSTCC | 1620 | Db |
| b | 1561 | CAACGGGGAGGTTGCTGAAAGTTCTCAAATGTCGCTGTAGCACTAGSTCC | 1620 | Qy |
| b | 1621 | TCTGTTCTTAATCTAATTACTTCTAGGTTAGCCAGTGCTCTCCACCTATGGGAGATG | 1680 | Db |
| b | 1621 | TCTGTTCTTAATCTAATTACTTCTAGGTTAGCCAGTGCTCTCCACCTATGGGAGATG | 1680 | Qy |
| b | 1680 | 1740 | Db | |
| b | 1740 | 1740 | Qy | |
| b | 1741 | GTTCGGTAACCTGGCACTTATCTAACTTAATACTCAAACATGAGCTT | 1800 | Db |
| b | 1741 | GTTCGGTAACCTGGCACTTATCTAACTTAATACTCAAACATGAGCTT | 1800 | Qy |
| b | 1801 | AGTTAACTGTTGACTTTCTCATCTGACCTCTGCTTATCCAGCCATT | 1860 | Db |
| b | 1801 | AGTTAACTGTTGACTTTCTCATCTGACCTCTGCTTATCCAGCCATT | 1860 | Qy |
| b | 1861 | TTGACAGTATATGGAAACCTCTCAACTGTCCTCTATGATCTAACCCCTT | 1920 | Db |
| b | 1861 | TTGACAGTATATGGAAACCTCTCAACTGTCCTCTATGATCTAACCCCTT | 1920 | Qy |
| b | 1920 | 1980 | Db | |
| b | 1920 | 1980 | Qy | |
| b | 1981 | CTATATAGTCCACAGGAGGTTATAACTCCATTAAAGTCTTGTGCTC | 2040 | Db |
| b | 1981 | CTATATAGTCCACAGGAGGTTATAACTCCATTAAAGTCTTGTGCTC | 2040 | Qy |
| b | 2041 | TGCCACCTTCAGGAACTCCAAATCTAGATTCTACTATTAACCTGTC | 2100 | Db |
| b | 2041 | TGCCACCTTCAGGAACTCCAAATCTAGATTCTACTATTAACCTGTC | 2100 | Qy |
| b | 2100 | 2200 | Db | |
| b | 2200 | 2200 | Qy | |
| b | 2201 | TCTAATTCTCTAGTCATGAGATCTACTCTCCAGGAGCTTATGCTACC | 2280 | Db |
| b | 2201 | TCTAATTCTCTAGTCATGAGATCTACTCTCCAGGAGCTTATGCTACC | 2280 | Qy |
| b | 2280 | 2280 | Db | |

| | | | | |
|----|------|--|------|----|
| Db | 2281 | CCTCCAAAGATGTCATGAGTCCCTTCATTCACTACATCAGCATCCATCACCCA | 2340 | Qy |
| Qy | 2341 | TGTGTGTTACTGATACTATGGCTCTCTGTTAGGACTAGTCACAGAGCC | 2400 | Db |
| Db | 2341 | TGTGTGTTACTGATACTATGGCTCTCTGTTAGGACTAGTCACAGAGC | 2400 | Qy |
| Qy | 2401 | TACATGGTGCTCTCTGTGATGTTAGTGTGTTCTGTTAGGAGTAGTCACAGAGC | 2460 | Db |
| Db | 2401 | TACATGGTGCTCTCTGTGATGTTAGTGTGTTCTGTTAGGAGTAGTCACAGAGC | 2460 | Qy |
| Qy | 2461 | GACTTAACTTTATGTAATGAAATAACCCATCTCATGAGTGTACTTTGTCGA | 2520 | Db |
| Db | 2461 | GACTTAACTTTATGTAATGAAATAACCCATCTCATGAGTGTACTTTGTCGA | 2520 | Qy |
| Qy | 2521 | AGACCCGGTTCTGAGCATTATTTATGATTATTCTTACCATGAG | 2580 | Db |
| Db | 2521 | AGACCCGGTTCTGAGCATTATTTATGATTATTCTTACCATGAG | 2580 | Qy |
| Qy | 2581 | GAGGTACTACATCTACATCTCTTATTATGATAAAGTAAGCCAGAGAATGATT | 2640 | Db |
| Db | 2581 | GAGGTACTACATCTACATCTCTTATTATGATAAAGTAAGCCAGAGAATGATT | 2640 | Qy |
| Qy | 2641 | AACCTACCCAAAGTATGAGTGTAGTGTAGCAGGSCCAAATTCAACCGTCCCCTA | 2700 | Db |
| Db | 2641 | AACCTACCCAAAGTATGAGTGTAGTGTAGCAGGSCCAAATTCAACCGTCCCCTA | 2700 | Qy |
| Qy | 2701 | TTACGTGTTAATCTGTCATGCTGCTCTGTCATCTATGGCTGAGATGAGAT | 2760 | Db |
| Db | 2701 | TTACGTGTTAATCTGTCATGCTGCTCTGTCATCTATGGCTGAGATGAGAT | 2760 | Qy |
| Qy | 2761 | CTGCCTCCGTAAGGAGGAAAGTATGGAGGAGATGGAGATGGAGAACGATTAAT | 2820 | Db |
| Db | 2761 | CTGCCTCCGTAAGGAGGAAAGTATGGAGGAGATGGAGATGGAGAACGATTAAT | 2820 | Qy |
| Qy | 2821 | CAGGGAAAGTCAACAGGAGCTGAGCTGAAAGTGTACTCTGTGTGTC | 2880 | Db |
| Db | 2821 | CAGGGAAAGTCAACAGGAGCTGAGCTGAAAGTGTACTCTGTGTGTC | 2880 | Qy |
| Qy | 2881 | GCATGACATCTGATGAACTGGTGTGACACACACATACATGGCTGATGTTACATA | 2940 | Db |
| Db | 2881 | GCATGACATCTGATGAACTGGTGTGACACACACATACATGGCTGATGTTACATA | 2940 | Qy |
| Qy | 2941 | TTTTTGTTAGTTGTGTTATTTATATATTTATGAAATGAAATGCTAAT | 3000 | Db |
| Db | 2941 | TTTTTGTTAGTTGTGTTATTTATATATTTATGAAATGAAATGCTAAT | 3000 | Qy |
| Qy | 3001 | TTCTAAGGCCAACATTGGCGGCAATTACATTGTCCTCTAACCTTGTG | 3060 | Db |
| Db | 3001 | TTCTAAGGCCAACATTGGCGGCAATTACATTGTCCTCTAACCTTGTG | 3060 | Qy |
| Qy | 3061 | AGATGATCAGATGGATTACTGGCCTGAGATGATATCTACATCTATCTATC | 3120 | Db |
| Db | 3061 | AGATGATCAGATGGATTACTGGCCTGAGATGATATCTACATCTATCTATC | 3120 | Qy |
| Qy | 3121 | TATATCTATCTATCTATCTATCTATCTATCTATCTATCTATCTATC | 3180 | Db |
| Db | 3121 | TATATCTATCTATCTATCTATCTATCTATCTATCTATCTATC | 3180 | Qy |
| Qy | 3181 | AAATATGTTGAAAGTTAAAGATTTCAGACTTTAGATCTGGATTGCAAAT | 3240 | Db |
| Db | 3181 | AAATATGTTGAAAGTTAAAGATTTCAGACTTTAGATCTGGATTGCAAAT | 3240 | Qy |
| Qy | 3241 | GTACCCCTTCTCTACATTAACCACTGTGAAACAATACATTTTATTCATC | 3300 | Db |
| Db | 3241 | GTACCCCTTCTCTACATTAACCACTGTGAAACAATACATTTTATTCATC | 3300 | Qy |
| Qy | 3301 | AAATGTTCTGACTCTGCTGATGAAACCGACACTGTCAGAACGCTTGGATA | 3360 | Db |
| Db | 3301 | AAATGTTCTGACTCTGCTGATGAAACCGACACTGTCAGAACGCTTGGATA | 3360 | Qy |
| Qy | 3361 | CATGCTGGCAAGCTATAGTTGCTCTCATTAACCTCTATTCAGTCCTCATCA | 3420 | Db |

COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/ms-DOS
 SOFTWARE: Patent Release #1.0, Version #1.25
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US108148, 058A
 FILING DATE: 04-Nov-1993
 CLASSIFICATION: 435
 ATTORNEY / AGENT INFORMATION:
 NAME: MOI, LESLIE A.
 REGISTRATION NUMBER: 37,047
 REFERENCE/DOCKET NUMBER: 028722-074
 TELECOMMUNICATION INFORMATION:
 TELEPHONE: 415-854-7400
 TELEFAX: 415-854-8275
 INFORMATION FOR SEQ ID NO: 12:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 2240 base pairs
 TYPE: nucleic acid
 STRANDEDNESS: double
 TOPOLOGY: linear
 US - 08-148-058A-12

Query Match 42.9%; Score 2240; DB 1; Length 2240;
 Best Local Similarity 100 %; Pred. No. 0;
 Mismatches 0; Indels 0; Gaps 0;
 Matches 2240; Conservative 0; Bimatches 0;

| Query | Subject |
|-------|--|
| QY | 1 GRATCTTGTAAATGGGCTTAGGGGTGGGGTTAACCCATTAGT |
| Db | 4861 GCTGCTTCTTCTTATGGCTCATTAACCTTAATTCCTGCACT |
| Qy | 4921 TAGGCAGACTGCTTGAATTCTGCTGTGAGGAAATTGTTATTC |
| Db | 4921 TAGGCAGACTGCTTGAATTCTGCTGTGAGGAAATTGTTATTC |
| Qy | 4981 AAATTAATAAGTTGCAAGTTTCTGCCAACAGACTCTGTCTTGAC |
| Db | 4981 AAATTAATAAGTTGCAAGTTTCTGCCAACAGACTCTGTCTTGAC |
| Qy | 5041 ATAAATCAAATACCGCTATGCTTAATTGGCAATGTCCTAACCTA |
| Db | 5041 ATAAATCAAATACCGCTATGCTTAATTGGCAATGTCCTAACCTA |
| Qy | 5101 GGAATACCAATAAGAACAGATAACACACAAAGGTACTAGTAA |
| Db | 5101 GGAATACCAATAAGAACAGATAACACACAAAGGTACTAGTAA |
| Qy | 5161 GAAAGAGTTAAAGAATTTCAGATGATTCCATATGTCCTCACCACGCA |
| Db | 5161 GAAAGAGTTAAAGAATTTCAGATGATTCCATATGTCCTCACCACGCA |
| Qy | 5221 AACN 5224 |
| Db | 5221 AACN 5224 |

RESULT 8
 US-08-148-058A-12
 Sequence 12, Application US/08148058A
 Patent No. 580407
 GENERAL INFORMATION:
 APPLICANT: TAMAKI, TAIKI
 APPLICANT: NAKABAYASHI, HIDEKAZU
 TITLE OF INVENTION: IMPROVED METHOD OF EXPRESSING GENES IN
 TITLE OF INVENTION: MAMMALIAN CELLS
 NUMBER OF SEQUENCES: 46
 CORRESPONDENCE ADDRESS:
 ADDRESSEE: BURNS, DOANE, SWECKER & MATHIS
 STREET: 699 PRINCE STREET
 CITY: ALEXANDRIA
 STATE: VA
 ZIP: 22313-1404
 COUNTRY: USA
 COMPUTER READABLE FORM:
 MEDIUM TYPE: Floppy disk

Qy 721 TAAGGGTGGTACATTTGTCAGCATTTGAGGAGAGTGAATTGAAAC 780
 Db 721 TAAAGGGTGGTACATTTGTCAGCATTTGAGGAGAGTGAATTGAAAC 780
 Qy 781 ATTTCACTAACACCCAACTCAGAACACAAATGAAAGATTCAGAAA 840
 Db 781 ATTTCACTAACACCCAACTCAGAACACAAATGAAAGATTCAGAAA 840
 Qy 841 GTGAGATAAGAGAGGAATTCTCACACCCACCGTATAGCTCAACTGCTGAGAA 900
 Db 841 GTGAGATAAGAGAGGAATTCTCACACCCACCGTATAGCTCAACTGCTGAGAA 900
 Qy 901 GTATATCTAATTACACTAACATCTGTAATAATGATAATTAGTCATT 960
 Db 901 GTATATCTAATTACACTAACATCTGTAATAATGATAATTAGTCATT 960
 Qy 961 TTAACTCTAATAGTACAGGATTTCACACCCACGTATAGCTCAACTGCTGAGAA 990
 Db 961 TTAACTCTAATAGTACAGGATTTCACACCCACGTATAGCTCAACTGCTGAGAA 990
 Qy 1021 ACTTGAGGGATAGTACATTCTGATGATAATGAGAAATTAAACAGTTGAAAT 1020
 Db 1021 ACTTGAGGGATAGTACATTCTGATGATAATGAGAAATTAAACAGTTGAAAT 1020
 Qy 1081 TTGCCTGTCATACAGTAAATTGACCATAGAGATATTCTGATGATAGTAAATAA 1080
 Db 1081 TTGCCTGTCATACAGTAAATTGACCATAGAGATATTCTGATGATAGTAAATAA 1080
 Qy 1141 CTTCTATACCAAACTTCAGTTACTGTTCCATGTTGCTTGAGTGGCTCACAGT 1200
 Db 1141 CTTCTATACCAAACTTCAGTTACTGTTCCATGTTGCTTGAGTGGCTCACAGT 1200
 Qy 1201 TATGAAAAAGAACGGATCAGAAATCATCATGCAAACCATCTGGAACACTGT 1260
 Db 1201 TATGAAAAAGAACGGATCAGAAATCATCATGCAAACCATCTGGAACACTGT 1260
 Qy 1261 ACTTAGCTACTAACTTGTGAGCTAAACACATAGATTGAGGAACTTGTGTGAGCT 1320
 Db 1261 ACTTAGCTACTAACTTGTGAGCTAAACACATAGATTGAGGAACTTGTGTGAGCT 1320
 Qy 1321 ACAAACTCTGTTCAAGCTCTTATTGTTGTTGAAATTGTTCTCAT 1380
 Db 1321 ACAAACTCTGTTCAAGCTCTTATTGTTGTTGAAATTGTTCTCAT 1380
 Qy 1381 GGTTCCTCTTCACTGCTATTTCTAACCTCACATGGCTACATACTGTC 1440
 Db 1381 GGTTCCTCTTCACTGCTATTTCTAACCTCACATGGCTACATACTGTC 1440
 Qy 1441 TCGAAAGTTATCTCCAAATCTCTGCTAACCTGTTGAGGATAAA 1500
 Db 1441 TCGAAAGTTATCTCCAAATCTCTGCTAACCTGTTGAGGATAAA 1500
 Qy 1501 AGTAGATTCAATGCCATGACATGACATGACATGACATGACATGACAT 1560
 Db 1501 AGTAGATTCAATGCCATGACATGACATGACATGACATGACATGACAT 1560
 Qy 1561 CAACACGGGAGTTGCTGGAGTTCTAAATGTGCTCTGAGACATGGTC 1620
 Db 1561 CAACACGGGAGTTGCTGGAGTTCTAAATGTGCTCTGAGACATGGTC 1620
 Qy 1620 CAACACGGGAGTTGCTGGAGTTCTAAATGTGCTCTGAGACATGGTC 1620
 Qy 1621 TCTTGTTCCTTAATCTAAATCTAAATCTAAATCTAAATCTAAATCTAA 1680
 Db 1621 TCTTGTTCCTTAATCTAAATCTAAATCTAAATCTAAATCTAAATCTAA 1680
 Qy 1681 AGGTGAAACGGAGCTGATGAAATTAATCACTAATGAGCTGAGACT 1740
 Db 1681 AGGTGAAACGGAGCTGATGAAATTAATCACTAATGAGCTGAGACT 1740
 Qy 1741 GTTGGGTTAACTGGTCACTTATCTAACTAATGAGCTGAGACT 1800
 Db 1741 GTTGGGTTAACTGGTCACTTATCTAACTAATGAGCTGAGACT 1800

Qy 1801 AGTTACTAAGCTTGTGTTCTCATACCCTCAGCTTATCAGGCCACTTAT 1860
 Db 1801 AGTACTAAGCTTGTGTTCTCATACCCTCAGCTTATCAGGCCACTTAT 1860
 Qy 1861 TTGACACTATATTGCGAAACTCTCTAACCTGCTGCTCTTCATAGCTTATCCCTT 1920
 Db 1861 TTGACACTATATTGCGAAACTCTCTAACCTGCTGCTCTTCATAGCTTATCCCTT 1920
 Qy 1921 TGAACCAAAGGAGCTTCAAAATCAATGTTTATAGTCCCTTTGTT 1980
 Db 1921 TGAACCAAAGGAGCTTCAAAATCAATGTTTATAGTCCCTTTGTT 1980
 Qy 1981 CTATAATAGTCCAGAGGGTATAACTCTGTTAAGTGTGAGTGTGCCCCT 2040
 Db 1981 CTATAATAGTCCAGAGGGTATAACTCTGTTAAGTGTGAGTGTGCCCCT 2040
 Qy 2041 TCCTAACATTGCGAGATTCCAATCTCTAGTTTCTACTTAACTTGTGCTC 2100
 Db 2041 TCCTAACATTGCGAGATTCCAATCTCTAGTTTCTACTTAACTTGTGCTC 2100
 Qy 2101 TTCAAACTGCAATTCTCATCCCTAAGTGTGCAATTGTTCCCTACGGTTGTT 2160
 Db 2101 TTCAAACTGCAATTCTCATCCCTAAGTGTGCAATTGTTCCCTACGGTTGTT 2160
 Qy 2161 TTCCACACCTTTGAGTTCTGGAACCTATACCCCTCCCTCATTTGCCACC 2220
 Db 2161 TTCCACACCTTTGAGTTCTGGAACCTATACCCCTCCCTCATTTGCCACC 2220
 Qy 2221 TCTAATTCTCTCAGACT 2240
 Db 2221 TCTAATTCTCTCAGACT 2240

RESULT 9
 US-08-478-042-12
 ; Sequence 12, Application US/08478042
 ; Patent No. 580738
 ; GENERAL INFORMATION:
 ; APPLICANT: TAMOKI, TAIKI
 ; ADDRESSER: NAKABAYASHI, HIDEKAZU
 ; TITLE OF INVENTION: IMPROVED METHOD OF EXPRESSING GENES IN
 ; NUMBER OF SEQUENCES: 46
 ; CORRESPONDENCE ADDRESS:
 ; ADDRESSER: BURNS, DOANE, SWECKER & MATHIS
 ; STREET: 699 PRINCE STREET
 ; CITY: ALEXANDRIA
 ; STATE: VA
 ; COUNTRY: USA
 ; ZIP: 22313-1404
 ; COMPUTER READABLE FORM:
 ; COMPUTER TYPE: Floppy disk
 ; COMPUTER: IBM PC compatible
 ; OPERATING SYSTEM: PC-DOS/MS-DOS
 ; SOFTWARE: Patient in Release #1.0, Version #1.25
 ; CURRENT APPLICATION DATA:
 ; APPLICATION NUMBER: US/08/478, 042
 ; FILING DATE: 07-JUN-1995
 ; CLASSIFICATION: 435
 ; PRIORITY APPLICATION DATA:
 ; APPLICATION NUMBER: US 08/148, 058
 ; FILING DATE: 04-NOV-1993
 ; ATTORNEY/AGENT INFORMATION:
 ; NAME: MOOL, LESLIE A.
 ; REGISTRATION NUMBER: 37, 047
 ; REFERENCE/DOCKET NUMBER: 028722-126
 ; TELECOMMUNICATION INFORMATION:
 ; TELEPHONE: 415-854-7100
 ; TELEX/FAX: 415-854-8275
 ; INFORMATION FOR SEQ ID NO: 12:
 ; LENGTH: 2240 base pairs
 ; TYPE: nucleic acid

;;
 STRANDBIENESS: double
 TOPOLOGY: linear
 US-08-478-042-12

Query Match 42.9%; Score 2240; DB 1; Length 2240;
 Best Local Similarity 100.0%; Pred. No. 0; Mismatches 0; Indels 0; Gaps 0;
 Matches 2240; Conservative 0; Mi mismatches 0;

Qy 1 GAATCTTAGAATATGGGTTAGCGGTGGGTAAATTCTGTTTACCCATAGT 60
 Db 1 GAATCTTAGAATATGGGTTAGCGGTGGGTAAATTCTGTTTACCCATAGT 60
 Qy 61 GAGATAGCATGGTTAATGGGTTAGGGGTGGTGAATTCTGTTTACCCATAGT 60
 Db 61 GAGATAGCATGGTTAATGGGTTAGGGGTGGTGAATTCTGTTTACCCATAGT 60
 Qy 121 AACAGACTATGGCTGGAGGCTTGAGGATCTGGATCTCAACACTGGTGTATC 180
 Db 121 AACAGACTATGGCTGGAGGCTTGAGGATCTGGATCTCAACACTGGTGTATC 180
 Qy 181 GTTCTATGGGCTTGTGTTAGCTGGTGCACAGACATCACATTGAAATTAGG 120
 Db 181 GTTCTATGGGCTTGTGTTAGCTGGTGCACAGACATCACATTGAAATTAGG 120
 Qy 121 AACAGACTATGGCTGGAGGCTTGAGGATCTGGATCTCAACACTGGTGTATC 180
 Db 121 AACAGACTATGGCTGGAGGCTTGAGGATCTGGATCTCAACACTGGTGTATC 180
 Qy 241 AAGCCCAAGGTACTGTCTCTTCATCTGTTGGGCTCTGGGCTTGAAATTCT 300
 Db 241 AAGCCCAAGGTACTGTCTCTTCATCTGTTGGGCTCTGGGCTTGAAATTCT 300
 Qy 301 GAGAAATAATAACATTCATAATGTTCTGGTGTAGATGAGATGAGATGTCAT 360
 Db 301 GAGAAATAATAACATTCATAATGTTCTGGTGTAGATGAGATGAGATGTCAT 360
 Qy 361 TCATTGTTATCATGATGAAATGAGAACATTAATCTGTTAGTACAATCT 420
 Db 361 TCATTGTTATCATGATGAAATGAGAACATTAATCTGTTAGTACAATCT 420
 Qy 421 GAGGGTAGGGCTGGTACTATCAATTCTTATTAAGAATRACTTATCTTATTA 480
 Db 421 GAGGGTAGGGCTGGTACTATCAATTCTTATTAAGAATRACTTATCTTATTA 480
 Qy 481 TGCCTGTGACAAATGTTGTCGGACACAGGAATCACAAAGATGAGCTTGGATT 540
 Db 481 TGCCTGTGACAAATGTTGTCGGACACAGGAATCACAAAGATGAGCTTGGATT 540
 Qy 541 AGAAAGTTAATGTCGAGGAAATTAATCACATGCTCAAACTGATGTTACCATCAA 600
 Db 541 AGAAAGTTAATGTCGAGGAAATTAATCACATGCTCAAACTGATGTTACCATCAA 600
 Qy 601 ACAAGAGGTTCATGAGAAATTAATGTCGAGGTTAATAGTTCTCAAGGTGAGGG 660
 Db 601 ACAAGAGGTTCATGAGAAATTAATGTCGAGGTTAATAGTTCTCAAGGTGAGGG 660
 Qy 661 CTCTCTCTAGTAGAGACTATCAGAAATCATCAGGAAATTATTGATAGACT 720
 Db 661 CTCTCTCTAGTAGAGACTATCAGAAATCATCAGGAAATTATTGATAGACT 720
 Qy 721 TAAGGGTTGGGACATTTGTCANGCATTTGGAAGAGGAGGATGAAAC 780
 Db 721 TAAGGGTTGGGACATTTGTCANGCATTTGGAAGAGGAGGATGAAAC 780
 Qy 781 ATTTCACACTACCAACCAACCAACCAACAAACAAAAAATGAAAGAAATCTGAAACA 840
 Db 781 ATTTCACACTACCAACCAACCAACCAACAAACAAAAAATGAAAGAAATCTGAAACA 840
 Qy 841 GTGAGAATAGAGAGGAAATTCTCACACACCACAGTATGCTCACTGCTGAGAA 900
 Db 841 GTGAGAATAGAGAGGAAATTCTCACACACCACAGTATGCTCACTGCTGAGAA 900
 Qy 901 GTATATCTATTTAACCTAACATCATGCTAATATGATTAATTTCTGTCATT 960
 Db 901 GTATATCTATTTAACCTAACATCATGCTAATATGATTAATTTCTGTCATT 960
 Qy 961 TTTAATGCTATAAGTACCAAGCATTAGANGATATTTCATTPATATCAAATAA 1020

Db 2041 TCCCAACTTGCAGGAATCCCATACTAGTATTCTACTATTAACATTGTC 2100
 Qy 2101 TTCAAACGTGATTCTCATTCCTAAGTGCACTGCAACAGGGTCAGTACTTC 2160
 Db 2101 TTCAAACGTGATTCTCATTCCTAAGTGCACTGCAACAGGGTCAGTACTTC 2160
 Qy 2161 TTCCACCACTTACATTCTGGAACACTATACCTCCCTCATTTGCCACC 2220
 Db 2161 TTCCACCACTTACATTCTGGAACACTATACCTCCCTCATTTGCCACC 2220
 Qy 2221 TCTAATTCTTCAAGACT 2240
 Db 2221 TCTAATTCTTCAAGACT 2240

RESULT 10
 US-08-645-215-12
 Sequence 12, Application US/08645215
 Patent No. 5827686
 GENERAL INFORMATION:
 APPLICANT: NAKABAYASHI, HIDEKAZU
 TITLE OF INVENTION: IMPROVED METHOD OF EXPRESSING GENES IN
 NUMBER OF SEQUENCES: 46
 CORRESPONDENCE ADDRESS:
 ADDRESSEE: BURNS, DOANE, SWCKER & MATHIS, L.L.P.
 STREET: 699 PRINCE STREET
 CITY: ALEXANDRIA
 STATE: VA
 COUNTRY: USA
 ZIP: 22313-1404
 COMPUTER READABLE FORM:
 MEDIUM TYPE: Floppy disk
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: Patent Release #1.0, Version #1.25
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/08/08645,215
 FILING DATE: 13-MAY-1996
 CLASSIFICATION: 435
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: US 08/148,058
 FILING DATE: 04-NOV-1993
 ATTORNEY / AGENT INFORMATION:
 NAME: MOI, LESLIE A.
 REGISTRATION NUMBER: 37,047
 REFERENCE/DOCKET NUMBER: 08872-135
 TELECOMMUNICATION INFORMATION:
 TELEPHONE: 415-854-7400
 TELEX/FAX: 415-854-8275
 INFORMATION FOR SEQ ID NO: 12:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 2240 base pairs
 TYPE: nucleic acid
 STRANDEDNESS: double
 TOPOLOGY: linear
 US-08-645-215-12

Query Match 42.9%; Score 2240; DB 1; length 2240;
 Best Local Similarity 100.0%; Pred. No. 0;
 Matches 2240; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 GAATCTTAGAAATATGGGGTAGGGGGTGGGGTGGTAATTCTGGTTACCCATAGT 60
 Db 1 GAATCTTAGAAATATGGGGTAGGGGGTGGGGTGGTAATTCTGGTTACCCATAGT 60
 Qy 61 GAGATAGCATGGGTTAATGCTTCACACACATCATTCTAGAATTAGG 120
 Db 61 GAGATAGCATGGGTTAATGCTTCACACACATCATTCTAGAATTAGG 120
 Qy 121 ACAGAGTATGGCTGGAGGCTTCAAGGAGCTTCAACTTGGTTGATCT 180
 Db 121 AACAGACTATGGCTGGAGCTTGGAGATCTGCTCATCACACTGGGTGATCT 180
 Qy 181 GTCTTATGGGCTGTTAAGCTGGCACTGCAACAGGGTCAGTACTTC 240
 Db 181 GTCTTATGGGCTGTTAAGCTGGCACTGCAACAGGGTCAGTACTTC 240
 Qy 241 AAGCCMAGGTACTGCTCTTCAATCTGTTGGGCTCTGGGCTGAATCT 300
 Db 241 AAGCCMAGGTACTGCTCTTCAATCTGTTGGGCTCTGGGCTGAATCT 300
 Qy 301 GAGAAATATAACATTCATAATGTCCTGGGAGATGAGTGGATGAGTGGCT 360
 Db 301 GAGAAATATAACATTCATAATGTCCTGGGAGATGAGTGGATGAGTGGCT 360
 Qy 361 TCATTGTTATCATGATGAGTGGACATTAGTGTATAACCTTAGTACACACT 420
 Db 361 TCATTGTTATCATGATGAGTGGACATTAGTGTATAACCTTAGTACACACT 420
 Qy 421 GAGGGTAGGGTGTGACTATTCAATTGTTGTCGGGACACGGATCACAAGT 480
 Db 421 GAGGGTAGGGTGTGACTATTCAATTGTTGTCGGGACACGGATCACAAGT 480
 Qy 481 TCCTGGACAAATGTTGTCGGGACACGGATCACAAGTGGCTTGAAATT 540
 Db 481 TCCTGGACAAATGTTGTCGGGACACGGATCACAAGTGGCTTGAAATT 540
 Qy 541 AAGAGTAAATGTCGGAAATAATCTGAAGTATCTGAGCTTACAAAGT 600
 Db 541 AAGAGTAAATGTCGGAAATAATCTGAAGTATCTGAGCTTACAAAGT 600
 Qy 601 ACAAGAGCTCATGAAATAATCTGAAGTTATAAGTGTCAAGGGGG 660
 Db 601 ACAAGAGCTCATGAAATAATCTGAAGTTATAAGTGTCAAGGGGG 660
 Qy 661 CTCTCTCTAGCTAGACTATCAGAAATCTGAAGGTTATAAGTGTCAAGGGGG 720
 Db 661 CTCTCTCTAGCTAGACTATCAGAAATCTGAAGGTTATAAGTGTCAAGGGGG 720
 Qy 721 TRAGGTGGGACATTGTCAGGAAATACTGAAGGTTATAAGTGTCAAGGGGG 780
 Db 721 TAAGGTGGGACATTGTCAGGAAATACTGAAGGTTATAAGTGTCAAGGGGG 780
 Qy 781 ATTCACCAACCAACCAATCCACAAACAAACAAAGGAATCTCAGAAC 840
 Db 781 ATTCACCAACCAACCAATCCACAAACAAACAAAGGAATCTCAGAAC 840
 Qy 841 GTGAGATAGAAAGAAATTCTCAACCCACAGGTAGCTAACCTGCTGAGAA 900
 Db 841 GTGAGATAGAAAGAAATTCTCAACCCACAGGTAGCTAACCTGCTGAGAA 900
 Qy 901 GTATATCTTAACTTAACTAAGCTAACATGCTAAATGTTAAATTACTGCTT 960
 Db 901 GTATATCTTAACTTAACTAAGCTAACATGCTAAATGTTAAATTACTGCTT 960
 Qy 961 TTAAATGCTTAACTAAGCTAACAGGCTTGGAGATTTCCATTATATCAAATAA 1020
 Db 961 TTAAATGCTTAACTAAGCTAACAGGCTTGGAGATTTCCATTATATCAAATAA 1020

Qy 1021 ACTTGAGGGTAGATGATTCATGATGAGTAAATTAACACAGATGAAATT 1080
 Db 1021 ACTTGAGGGTAGATGATTCATGATGAGTAAATTAACACAGATGAAATT 1080
 Qy 1081 TTGCTTGTCTACACATTAATGACATAGCATAGATTAAATTGTTGAAT 1140
 Db 1081 TTGCTTGTCTACACATTAATGACATAGCATAGATTAAATTGTTGAAT 1140
 Qy 1141 CTTCTATACACAGTCACTGTTGTCATGTTGCTCTGAGTGGCTCACAGACT 1200
 Db 1141 CTTCTATACACAGTCACTGTTGTCATGTTGCTCTGAGTGGCTCACAGACT 1200
 Qy 1201 TATGAAAGTAAACGGAACTCAAGATTCACATGCAAAAGCATGCTGAACTCTGT 1260
 Db 1201 TATGAAAGTAAACGGAACTCAAGATTCACATGCAAAAGCATGCTGAACTCTGT 1260

Patent No. 5843776
 GENERAL INFORMATION:
 APPLICANT: TAMAOKI, TAIKI
 APPLICANT: NAKABAYASHI, HIDEKAZU
 TITLE OF INVENTION: IMPROVED METHOD OF EXPRESSING GENES IN
 MAMMALIAN CELLS
 NUMBER OF SEQUENCES: 46
 CORRESPONDENCE ADDRESS:
 ADDRESSEE: BURNS, DONNE, SWECKER & MATHIS
 STREET: 699 PRINCE STREET
 CITY: ALEXANDRIA
 STATE: VA
 COUNTRY: USA
 ZIP: 22313-1404
 COMPUTER READABLE FORM:
 MEDIUM TYPE: FLOPPY disk
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: Patent Release #1.0, Version #1.25
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/08/466,604
 FILING DATE: 06-JUN-1995
 CLASSIFICATION: 435
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: US 08/148,058
 FILING DATE: 04-NOV-1993
 ATTORNEY/AGENT INFORMATION:
 NAME: MOOI, LESLIE A.
 REGISTRATION NUMBER: 37,047
 REFERENCE/DOCKET NUMBER: 028722-125
 TELECOMMUNICATION INFORMATION:
 TELEPHONE: 415-854-7400
 TELEFAX: 415-854-8275
 INFORMATION FOR SEQ ID NO: 12:
 LENGTH: 2240
 SEQUENCE CHARACTERISTICS:
 TYPE: nucleic acid
 STRANDEDNESS: double
 TOPOLOGY: linear
 US-08-466-604-12

Query Match 42.9%; Score 2240; DB 2; Length 2240;
 Best Local Similarity 100.0%; Pred. No. 0;
 Matches 2240; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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| QY | 1 | GAATTCTTAAATGAGGTAGGGTAAATGCTGTTCACCCATAGGT | 60 |
| Db | 61 | GAGATAGCATGGGTTAATGCTTCAACACACATCATTAGAATTAGG | 120 |
| QY | 121 | ACAGACATATGGGTAGGGTAGGGTAACTTGCTCTCTACATAACACTCGGTCTATCT | 180 |
| Db | 121 | ACAGACATATGGGTAGGGTAGGGTAACTTGCTCTCTACATAACACTCGGTCTATCT | 180 |
| QY | 181 | GTTCTATGGGACTGTTAACGTTAGCTGCACTGTTACGAGGTGTT | 240 |
| Db | 181 | GTCTATGGGACTGTTAACGTTAGCTGCACTGTTACGAGGTGTT | 240 |
| QY | 241 | AAGCCAGGACTGCTCTTCATCTGTTGGGCCCTGGGCTGATMCT | 300 |
| Db | 241 | AAGCCAGGACTGCTCTTCATCTGTTGGGCCCTGGGCTGATMCT | 300 |
| QY | 301 | GAGAAATAACATTCATAATGTCCTGTTGAGATGAGATGAGATGTCAT | 360 |
| Db | 301 | GAGAAATAACATTCATAATGTCCTGTTGAGATGAGATGAGATGTCAT | 360 |
| QY | 361 | TCTATGTTCAATGATGAGGAACTAGTGTAACTCTTACACATCT | 420 |
| Db | 361 | TCTATGTTCAATGATGAGGAACTAGTGTAACTCTTACACATCT | 420 |

RESULT 11
 ; Sequence 12, Application US/08466604
 ; Sequence 12, Application US/08466604

QY 421 GAGGGTAGGGTGTACTATTCAATTCTTATAAAGATACCTATTCTTTATTAA 480
 Db 421 GAGGGTAGGGTGTACTATTCAATTCTTATAAAGATACCTATTCTTTATTAA 480
 QY 481 TCGTGTGACAATGTTGTTGGGGACACAGGAATCACAAAGATGCTTGAAATT 540
 Db 481 TCGTGTGACAATGTTGTTGGGGACACAGGAATCACAAAGATGCTTGAAATT 540
 QY 541 ANGAAGTAAATGTCAGGAAATTACATAGTTACAGGAACTATGATACCATCAA 600
 Db 541 ANGAAGTAAATGTCAGGAAATTACATAGTTACAGGAACTATGATACCATCAA 600
 QY 601 ACAAGAGGTCCATGAGAAATTCTGAGAGTTATACTGTCAAGGTGAGAGGG 660
 Db 601 ACAAGAGGTCCATGAGAAATTCTGAGAGTTATACTGTCAAGGTGAGAGGG 660
 QY 661 CTCCTCTCTACTGTAGAGACTATCGAAATACTTCAGGGATAATTTRATAGACT 720
 Db 661 CTCCTCTCTACTGTAGAGACTATCGAAATACTTCAGGGATAATTTRATAGACT 720
 QY 721 TAGGGTGGGACATTGTCAGGATTGAGAGGATGGAGAGAGGATTTGAAAC 780
 Db 721 TAGGGTGGGACATTGTCAGGATTGAGAGGATGGAGAGAGGATTTGAAAC 780
 QY 781 ATTTCACTAACCAACCACATCCACAAACAAATACTGAGAACTGAGGAGG 840
 Db 781 ATTTCACTAACCAACCACACCCGAACTCCACAAACAAATACTGAGAAAC 840
 QY 841 GTGAGATAAGAGAGGATTCTCAACACCCACACTATGCTCAACTGCTGAGAA 900
 Db 841 GTGAGATAAGAGAGGATTCTCAACACCCACACTATGCTCAACTGCTGAGAA 900
 QY 901 GTATATCTCTATTTAACATCATCTTAATGATAATTTCTGATTGCTT 960
 Db 901 GTATATCTCTATTTAACATCATCTTAATGATAATTTCTGATTGCTT 960
 QY 961 TTAAATGCTATAAGTACACGCAATTAGGATTTCTGATTGCTT 1020
 Db 961 TTAAATGCTATAAGTACACGCAATTAGGATTTCTGATTGCTT 1020
 QY 1021 ACTTGAGGGATAGATCATTCTCATGATATAGGAAATAAAACAGTTGAAATT 1080
 Db 1021 ACTTGAGGGATAGATCATTCTCATGATATAGGAAATAAAACAGTTGAAATT 1080
 QY 1081 TTGCCCTGTCATACAGCTAATTATGACCATAGACATTAAATTGTTGAAT 1140
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 Db 1141 CTTCATACACGAACTTCAACTTCAATGAGCTTCAACGACT 1200
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 Db 1201 TATGAAAAAGAACGAACTTCAATGAGCTTCAACGACT 1260
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 Db 1261 ACTTAGGACTAACTTGAGGAACTACACAGATAGTGAGGATGTTGTTGTTAGAT 1320
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 Db 1441 TCGAAGCTTATGATTCGAATATCTATGCTCAACTTGTGTTGGAAATTGCTGTTCTCAT 1500
 QY 1501 AGTACTTCAATGCACTCAACTTCCACTTGGGGCTTAAGACGTTCAACATA 1560

RESULT 12
 US-09-949-016-1702
 Sequence 1702, Application US/09949016
 - Patent No. 6812339
 GENERAL INFORMATION:
 - APPLICANT: VENTER, J. Craig et al.
 - TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
 - TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
 - FILE REFERENCE: C1001307
 - CURRENT APPLICATION NUMBER: US/09/949, 016
 - CURRENT FILING DATE: 2000-04-14
 - PRIOR APPLICATION NUMBER: 6/0241, 755
 - PRIOR FILING DATE: 2000-10-20
 - PRIOR APPLICATION NUMBER: 6/0237, 768
 - PRIOR FILING DATE: 2000-10-03
 - PRIOR APPLICATION NUMBER: 6/0231, 498
 - PRIOR FILING DATE: 2000-03-08
 - NUMBER OF SEQ ID NOS: 20712
 SOFTWARE: FastSEQ for Windows Version 4.0
 SEQ ID NO: 1702
 LENGTH: 23497

US-09-033-428-1
; Sequence 1, Application US/09033428
; Patent No. 6354862
; GENERAL INFORMATION:
; APPLICANT: Little, Andrew
; APPLICANT: Lamparski, Henry
; APPLICANT: Schuur, Eric
; APPLICANT: Henderson, Daniel
; TITLE OF INVENTION: ADENOVIRUS VECTORS SPECIFIC FOR CELLS
; NUMBER OF SEQUENCES: 23
; CORRESPONDENCE ADDRESS:
; ADDRESS: MORRISON & FOERSTER
; STREET: 755 PAGE MILL ROAD
; CITY: PALO ALTO
; STATE: CA
; COUNTRY: USA
; ZIP: 94304-1018
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MC-DOS
; SOFTWARE: Patentin Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/09/033,428
; FILING DATE:
; CLASSIFICATION:
; ATTORNEY/AGENT INFORMATION:
; NAME: POLIZZI, CATHERINE M.
; REGISTRATION NUMBER: 40,130
; REFERENCE/DOCKET NUMBER: 34802-30004.00
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (415) 813-5600
; TELEFAX: (415) 494-0792
; TELEX: 706141 MRNFOERS SPO
; INFORMATION FOR SEQ ID NO: 1:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 822 base pairs
; TYPE: nucleic acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; US-09-033-428-1

Query Match 11.8%; Score 619; DB 3; Length 822;
Best Local Similarity 100.0%; Pred. No. 9.7e-137;
Matches 619; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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Db 1 GCATTCGTGCTGTTAGGATAGAACCTCTGGTCAAGCTCTCTTTATGGTTGCTTGG 60
Qy 1302 GATGTTGTTGCTGTTAGGATAGAACCTCTGGTCAAGCTCTCTTTATGGTTGCTTGG 1361
Db 61 GATGTTGTTGCTGTTAGGATAGAACCTCTGGTCAAGCTCTCTTTATGGTTGCTTGG 120
Qy 1362 AAATTGTCGTTCTCATGGTTCTCTTCACTGTTATTTCTACCACTCA 1421
Db 121 AAATTGTCGTTCTCATGGTTCTCTTCACTGTTATTTCTACCACTCA 180
Qy 1422 CATGGCTCACATACTGTCGCAAGCTTATCCAAATCTATCTCGGCCAAT 1481
Db 181 CATGGCTCACATACTGTCGCAAGCTTATCCAAATCTATCTCGGCCAAT 240
Qy 1482 CTGGTCCAGAGATAAAGTAGTATTCAAGCACATCACTGTCGCAAGCTTGGGGC 1541
Db 241 CTGGTCCAGAGATAAAGTAGTATTCAAGCACATCACTGTCGCAAGCTTGGGGC 300
Qy 1542 TTAAGACGTTCAACATACAAACCGGGAGTTGCCTGAAATGTTCTCAAATGTT 1601
Db 301 TTAAGACGTTCAACATACAAACCGGGAGTTGCCTGAAATGTTCTCAAATGTT 360
Qy 1602 CCTGTGACACTAGTGGCTCTTGTCCTTAATCTATTTAGGCCAGTGCTCA 1661

RESULT 14
US-09-033-556-6
; Sequence 6, Application US/09033556
; Patent No. 6332700
; GENERAL INFORMATION:
; APPLICANT: Henderson, Daniel R.
; APPLICANT: Yu De Chao
; TITLE OF INVENTION: ADENOVIRUS VECTORS CONTAINING
; TITLE OF INVENTION: HETEROLOGOUS TRANSCRIPTION REGULATORY ELEMENTS AND METHODS
; TITLE OF INVENTION: OF USING SAME
; NUMBER OF SEQUENCES: 41
; CORRESPONDENCE ADDRESS:
; ADDRESSE: MORRISON & FOERSTER
; STREET: 755 PAGE MILL ROAD
; CITY: Palo Alto
; STATE: CA
; COUNTRY: USA
; ZIP: 94304-1018
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Diskette
; COMPUTER: IBM Compatible
; OPERATING SYSTEM: Windows
; SOFTWARE: PatisSEQ for Windows Version 2.0b
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/09/033,556
; FILING DATE:
; CLASSIFICATION:
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER:
; FILING DATE:
; ATTORNEY/AGENT INFORMATION:
; NAME: Catherine, Polizzi M
; REGISTRATION NUMBER: 40,130
; REFERENCE/DOCKET NUMBER: 34802-20010.00
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 650-813-5600
; TELEFAX: 650-494-0792
; TELEX: 706141
; INFORMATION FOR SEQ ID NO: 6:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 822 base pairs
; TYPE: nucleic acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; US-09-033-556-6

Query Match 11.8%; Score 619; DB 3; Length 822;
Best Local Similarity 100.0%; Pred. No. 9.7e-137;
Matches 619; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1242 GCATTCGTGCTGTTAGGATAGAACCTCTGGTCAAGCTCTCTTTATGGTTGCTTGG 1301
Db 1 GCATTCGTGCTGTTAGGATAGAACCTCTGGTCAAGCTCTCTTTATGGTTGCTTGG 60

GenCore version 5.1.6
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OM nucleic - nucleic search, using SW model

Run on: August 6, 2005, 22:19:03 ; Search time 492.167 Seconds
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Sequence: 1 GCATTCGCTGAACTCTGTAA.....TTCCACCACTGCCATTAACA 822

Scoring table: IDENTITY.NUC
Gapop 10.0 , Gapext 1.0

Searched: 7297361 seqB, 324162794 residues

Total number of hits satisfying chosen parameters: 14594722

Minimum DB seq length: 0
Maximum DB seq length: 200000000

Post-processing: Minimum Match 0%
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Listing first 45 summaries

Database : Published Applications NA:*

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21: /cgns_6/ptodata/2/pubpna/us10I_PUBCOMB.seq:/*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

| Result No. | Score | Query Match Length | DB ID | Description |
|------------|-------|--------------------|-------|--|
| 1 | 822 | 100.0 | 822 | 9 US-09-898-893-1 Sequence 1, Appli |
| 2 | 822 | 100.0 | 822 | 10 US-09-151-776-6 Sequence 6, Appli |
| 3 | 822 | 100.0 | 822 | 10 US-09-151-776-4 Sequence 44, Appli |
| 4 | 822 | 100.0 | 822 | 10 US-09-814-357-8 Sequence 8, Appli |
| 5 | 822 | 100.0 | 822 | 10 US-09-814-351-8 Sequence 14, Appli |
| 6 | 822 | 100.0 | 822 | 15 US-10-226-820-14 Sequence 6, Appli |
| 7 | 822 | 100.0 | 822 | 16 US-10-139-883-2 Sequence 6, Appli |
| 8 | 822 | 100.0 | 822 | 16 US-10-139-889-4 Sequence 4, Appli |
| 9 | 822 | 100.0 | 822 | 16 US-10-822-873-5 Sequence 5, Appli |
| 10 | 822 | 100.0 | 822 | 20 US-10-822-873-4 Sequence 44, Appli |
| 11 | 822 | 100.0 | 822 | 22 US-10-691-045-8 Sequence 8, Appli |
| 12 | 619 | 75.3 | 2240 | 10 US-09-828-455-12 Sequence 12, Appli |
| 13 | 619 | 75.3 | 5224 | 9 US-09-888-883-2 Sequence 2, Appli |
| 14 | 619 | 75.3 | 5224 | 10 US-09-151-376-5 Sequence 5, Appli |
| 15 | 619 | 75.3 | 5224 | 10 US-09-151-376-4 Sequence 45, Appli |
| 16 | 619 | 75.3 | 5224 | 10 US-09-814-357-16 Sequence 16, Appli |
| 17 | 619 | 75.3 | 5224 | 10 US-09-814-351-16 Sequence 15, Appli |
| 18 | 619 | 75.3 | 5224 | 15 US-10-226-820-15 Sequence 5, Appli |
| 19 | 619 | 75.3 | 5224 | 16 US-10-139-889-5 Sequence 45, Appli |
| 20 | 619 | 75.3 | 5224 | 16 US-10-139-889-45 Sequence 5, Appli |
| 21 | 619 | 75.3 | 5224 | 20 US-10-822-873-5 Sequence 45, Appli |
| 22 | 619 | 75.3 | 5224 | 20 US-10-822-873-45 Sequence 16, Appli |
| 23 | 619 | 75.3 | 5224 | 22 US-10-691-045-16 Sequence 58, Appli |
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| 25 | 411.2 | 50.0 | 5945 | 15 US-10-311-455-57 Sequence 60, Appli |
| 26 | 411.2 | 50.0 | 7001 | 15 US-10-72-086-60 Sequence 2, Appli |
| 27 | 157.6 | 19.2 | 7001 | 19 US-10-311-507-2 Sequence 60, Appli |
| 28 | 157.6 | 19.2 | 7001 | 20 US-10-480-846-60 Sequence 59, Appli |
| 29 | 139.8 | 17.0 | 7001 | 15 US-10-172-086-59 Sequence 1, Appli |
| 30 | 139.8 | 17.0 | 7001 | 19 US-10-311-507-1 Sequence 59, Appli |
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| 32 | 89.8 | 10.9 | 10415 | 21 US-10-703-224-2 Sequence 3, Appli |
| 33 | 73 | 8.9 | 228 | 9 US-09-947-925A-3 Sequence 20, Appli |
| 34 | 50 | 6.1 | 50 | 10 US-09-993-346-20 Sequence 352, App |
| 35 | 50 | 6.1 | 50 | 10 US-09-993-346-32 Sequence 7042, Ap |
| 36 | 49 | 6.0 | 81428 | 20 US-10-179-993-7042 Sequence 20740, Sequence 20771, Sequence 207471, Sequence 362, App |
| 37 | 48.6 | 5.9 | 616 | 17 US-10-027-632-207470 Sequence 207470, Sequence 207471, Sequence 207471, Sequence 362, App |
| 38 | 48.6 | 5.9 | 1137 | 13 US-10-027-632-207471 Sequence 207470, Sequence 207471, Sequence 362, App |
| 39 | 48.6 | 5.9 | 1137 | 17 US-10-027-632-207471 Sequence 207470, Sequence 207471, Sequence 362, App |
| 40 | 48.6 | 5.9 | 1137 | 17 US-10-027-632-207471 Sequence 207470, Sequence 207471, Sequence 362, App |
| 41 | 47.6 | 5.8 | 2750 | 10 US-09-854-867-362 Sequence 207470, Sequence 207471, Sequence 362, App |
| 42 | 47.6 | 5.8 | 2750 | 21 US-10-786-970A-362 Sequence 207470, Sequence 207471, Sequence 362, App |
| 43 | 43.8 | 5.3 | 5781 | 17 US-10-264-237-2827 Sequence 207470, Sequence 207471, Sequence 362, App |
| 44 | 43.8 | 5.3 | 23865 | 19 US-10-317-478-4 Sequence 207470, Sequence 207471, Sequence 362, App |
| 45 | 43.6 | 5.3 | 44030 | 11 US-09-997-722-172 Sequence 207470, Sequence 207471, Sequence 362, App |

ALIGNMENTS

RESULT 1
US-09-898-893-1
; Sequence 1, Application US/09898883
; Patent No. US20020164799A1
GENERAL INFORMATION:
APPLICANT: Little, Andrew
Lamparski, Henry
Schurz, Eric C
Henderson, Daniel
TITLE OF INVENTION: ADENOVIRUS VECTORS SPECIFIC FOR CELLS EXPRESSING ALPHA-FETOPROTEIN AND METHODS OF USE THEREOF
NUMBER OF SEQUENCES: 23
CORRESPONDENCE ADDRESS:
ADRESSEE: MORRISON & FOERSTER
STREET: 755 PAGE MILL ROAD
CITY: PALO ALTO
STATE: CA
COUNTRY: USA
ZIP: 94304-1018
COMPUTER READABLE FORM:
MEDIUM TYPE: floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patent Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/898,883
FILING DATE: 05-JUL-2001
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/033,428

FILING DATE: <Unknown>
ATTORNEY/AGENT INFORMATION:
NAME: POLIZZI, CATHERINE M.
REGISTRATION NUMBER: 40,130
REFERENCE/DOCKET NUMBER: 34802-30004.00

TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 813-5600
TELEFAX: (415) 494-0792
TELEX: 706341 MFSNFORS SFO

INFORMATION FOR SEQ ID NO: 1:

SEQUENCE CHARACTERISTICS:
LENGTH: 822 base pairs
TYPE: nucleic acid
STRANDBNESS: single
TOPOLY: linear
SEQUENCE DESCRIPTION: SEQ ID NO: 1:
US-09-898-883-1

Query Match 100.0%; Score 822; DB 9; Length 822;
Best Local Similarity 100.0%; Pred. No. 1.1e-216; Matches 822; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 GCATGGCTGAGACTCTGTACTTAGGACTAACTTGAGCATAACACATAGATTGAG 60
Db 1 GCATGGCTGAGACTCTGTACTTAGGACTAACTTGAGCATAACACATAGATTGAG 60
QY 61 GATTTGTTGCTTGTAGCATACAACTCTGGTCAAGCTCCCTTATTGCTGTCCTGG 120
Db 61 GATTTGTTGCTTGTAGCATACAACTCTGGTCAAGCTCCCTTATTGCTGTCCTGG 120
QY 121 AAATTGTTGCTTGTAGCATACAACTCTGGTCAAGCTCCCTTATTGCTGTCACCTCA 180
Db 121 AAATTGTTGCTTGTAGCATACAACTCTGGTCAAGCTCCCTTATTGCTGTCACCTCA 180
QY 181 CATGGCTACAATAACTCTGTGCAAGCTTATGCCAATATCTAACCTCTGAGCTCAAT 240
Db 181 CATGGCTACAATAACTCTGTGCAAGCTTATGCCAATATCTAACCTCTGAGCTCAAT 240
QY 241 CTGTTTCAGAGATAAAAGCTAGTATCAATGCACATCACAGCTGTCCTGGGGGG 300
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Db 301 TTAAGACAGTTCAACATACAAACCGGGAGTTCTGCTTAATCTAACTTGTGAGCTCA 360
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Db 421 TCCACCTATGGGAGATGAGCTGAAAGGAGGCCCTGATTATAATCTACTAAGTCAA 480
QY 481 TAGCTAGACCCAGGACTGTGCTGAATCTAACTAAATATC 540
Db 481 TAGCTAGACCCAGGACTGTGCTGAATCTAACTAAATATC 540
QY 541 CAAACTGACATGACTTACTAGTAACTCTGAGCTTGTGAGCTTGTGAGCTCA 600
Db 541 CAAACTGACATGACTTACTAGTAACTCTGAGCTTGTGAGCTTGTGAGCTCA 600
QY 601 CCTTATCGGCOACTATGAGCTGTCCTGACATAATAACAAATACCGCTAT 660
Db 601 CCTTATCGGCOACTATGAGCTGTCCTGACATAATAACAAATACCGCTAT 660
QY 661 GCTGTTPAATTGTGCAATTCGCCATTTCACCTTAACTGAGAAATCCATAGTACAGA 720
Db 661 GCTGTTPAATTGTGCAATTCGCCATTTCACCTTAACTGAGAAATCCATAGTACAGA 720
QY 721 TATACCAACAAAGGTACTGTGTTACAGGATTGCTGAAAGGATTAAGATTG 780
Db 721 TATACCAACAAAGGTACTGTGTTACAGGATTGCTGAAAGGATTAAGATTG 780

RESULT 2
US-09-151-376-6
Sequence 6, Application US/09151376
; Publication No. US20030044383A1
GENERAL INFORMATION:
APPLICANT: Henderson, D.R.
TITLE OF INVENTION: TISSUE SPECIFIC VIRAL VECTORS
FILE REFERENCE: 34802200021
CURRENT APPLICATION NUMBER: US/09/151, 376
CURRENT FILING DATE: 1998-09-10
EARLIER APPLICATION NUMBER: 08/669, 753
EARLIER FILING DATE: 1996-06-26
EARLIER APPLICATION NUMBER: 08/495, 034
NUMBER OF SEQ ID NOS: 71
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO: 6
LENGTH: 822
TYPE: DNA
ORGANISM: Homo sapiens
US-09-151-376-6

Query Match 100.0%; Score 822; DB 10; Length 822;
Best Local Similarity 100.0%; Pred. No. 1.1e-216; Matches 822; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 GCATGGCTGAGACTCTGTACTTAGGACTAACTTGAGCATAACACATAGATTGAG 60
Db 1 GCATGGCTGAGACTCTGTACTTAGGACTAACTTGAGCATAACACATAGATTGAG 60
QY 61 GATTTGTTGCTTGTAGCATACAACTCTGGTCAAGCTCCCTTATTGCTGTCCTGG 120
Db 61 GATTTGTTGCTTGTAGCATACAACTCTGGTCAAGCTCCCTTATTGCTGTCCTGG 120
QY 121 AAATTGTTGCTTGTAGCATACAACTCTGGTCAAGCTCCCTTATTGCTGTCACCTCA 180
Db 121 AAATTGTTGCTTGTAGCATACAACTCTGGTCAAGCTCCCTTATTGCTGTCACCTCA 180
QY 181 CATGGCTACAATAACTCTGTGCAAGCTTATGCCAATATCTAACCTCTGAGCTCAAT 240
Db 181 CATGGCTACAATAACTCTGTGCAAGCTTATGCCAATATCTAACCTCTGAGCTCAAT 240
QY 241 CTGTTTCAGAGATAAAAGCTAGTATCAATGCACATCACAGCTGTCCTGGGGGG 300
Db 241 CTGTTTCAGAGATAAAAGCTAGTATCAATGCACATCACAGCTGTCCTGGGGGG 300
QY 301 TTAAGACAGTTCAACATACAAACCGGGAGTTCTGCTTAATCTAACTTGTGAGCTCA 360
Db 301 TTAAGACAGTTCAACATACAAACCGGGAGTTCTGCTTAATCTAACTTGTGAGCTCA 360
QY 361 CCTGTAACATAAGGCTCTTGTGCTTAATCTAACTTGTGAGCTCA 420
Db 361 CCTGTAACATAAGGCTCTTGTGCTTAATCTAACTTGTGAGCTCA 420
QY 421 TCCACCTATGGGAGATGAGCTGAAAGGAGGCCCTGATTATAATCTACTAAGTCAA 480
Db 421 TCCACCTATGGGAGATGAGCTGAAAGGAGGCCCTGATTATAATCTACTAAGTCAA 480
QY 481 TAGCTAGACCCAGGACTGTGCTGAATCTAACTAAATATC 540
Db 481 TAGCTAGACCCAGGACTGTGCTGAATCTAACTAAATATC 540
QY 541 CAAACTGACATGACTTACTAGTAACTCTGAGCTTGTGAGCTCA 600
Db 541 CAAACTGACATGACTTACTAGTAACTCTGAGCTTGTGAGCTCA 600

QY 781 AGCATGATTCCATATGTCCTCCACCACTGCAAATAACA 822
Db 781 AGCATGATTCCATATGTCCTCCACCACTGCAAATAACA 822

RESULT 5
US-09-814-351-8

; Sequence 8, Application US/09814351

; Publication No. US2003014850A1

; GENERAL INFORMATION:

; APPLICANT: Yu, De-Chao

; APPLICANT: Li, Yuanhao

; APPLICANT: Henderson, Daniel R.

TITLE OF INVENTION: CELL-SPECIFIC ADENOVIRUS VECTORS
TITLE OF INVENTION: COMPRISING AN INTERNAL RIBOSOME ENTRY SITE
FILE REFERENCE: 348022001700
CURRENT APPLICATION NUMBER: US/09/814, 351
CURRENT FILING DATE: 2001-03-21
PRIOR FILING DATE: 2000-03-24
NUMBER OF SEQ ID NOS: 35

; SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO: 8
LENGTH: 822

TYPE: DNA

ORGANISM: Artificial Sequence

FEATURE: OTHER INFORMATION: AFR-TRE

US-09-814-351-8

Query Match 100.0%; Score 822; DB 10; Length 822;
Best Local Similarity 100.0%; Pred. No. 1.1e-216; Number of Mismatches 0; Indels 0; Gaps 0;

RESULT 6
US-10-226-820-14

; Sequence 14, Application US/10226820

; Publication No. US2003011955A1

; GENERAL INFORMATION:

; APPLICANT: Henderson, Daniel R.

; APPLICANT: Yu, De Chao

TITLE OF INVENTION: TARGET CELL-SPECIFIC ADENOVIRAL VECTORS
TITLE OF INVENTION: CONTAINING E3 AND METHODS OF USE THEREOF
FILE REFERENCE: 348022001300
CURRENT APPLICATION NUMBER: US/10/226, 820
CURRENT FILING DATE: 2002-08-21
PRIOR FILING DATE: 1999-12-29
PRIOR APPLICATION NUMBER: 60/114, 262

; PRIOR FILING DATE: 1998-12-30

RESULT 5
US-09-814-351-8

; Sequence 8, Application US/09814351

; Publication No. US2003014850A1

; GENERAL INFORMATION:

; APPLICANT: Yu, De-Chao

; APPLICANT: Li, Yuanhao

; APPLICANT: Henderson, Daniel R.

TITLE OF INVENTION: CELL-SPECIFIC ADENOVIRUS VECTORS
TITLE OF INVENTION: COMPRISING AN INTERNAL RIBOSOME ENTRY SITE
FILE REFERENCE: 348022001700
CURRENT APPLICATION NUMBER: US/09/814, 351
CURRENT FILING DATE: 2001-03-21
PRIOR FILING DATE: 2000-03-24
NUMBER OF SEQ ID NOS: 35

; SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO: 8
LENGTH: 822

TYPE: DNA

ORGANISM: Artificial Sequence

FEATURE: OTHER INFORMATION: AFR-TRE

US-09-814-351-8

Query Match 100.0%; Score 822; DB 10; Length 822;
Best Local Similarity 100.0%; Pred. No. 1.1e-216; Number of Mismatches 0; Indels 0; Gaps 0;

Page 5

RESULT 8
US-10-139-089-44

; Sequence 44, Application US/10139089

; Publication No. US2003015253A1

; GENERAL INFORMATION:

; APPLICANT: Henderson, D.R.

; APPLICANT: Schuur, E.R.

TITLE OF INVENTION: TISSUE SPECIFIC VIRAL VECTORS

FILE REFERENCE: 348024000221

CURRENT APPLICATION NUMBER: US/10/139,089

CURRENT FILING DATE: 2002-05-02

PRIOR APPLICATION NUMBER: 08/663,753

PRIOR FILING DATE: 1996-06-26

PRIOR APPLICATION NUMBER: 08/495,034

PRIOR FILING DATE: 1995-06-27

PRIOR APPLICATION NUMBER: 09/509,591

PRIOR FILING DATE: 2000-06-02

PRIOR APPLICATION NUMBER: 09/151,376

PRIOR FILING DATE: 1998-09-10

PRIOR APPLICATION NUMBER: 09/033,428

PRIOR FILING DATE: 1998-03-02

PRIOR APPLICATION NUMBER: 60/039,597

PRIOR FILING DATE: 1997-03-03

PRIOR APPLICATION NUMBER: 09/033,555

PRIOR FILING DATE: 1998-03-02

PRIOR APPLICATION NUMBER: 60/039,763

PRIOR FILING DATE: 1997-03-03

PRIOR APPLICATION NUMBER: 09/033,333

PRIOR FILING DATE: 1998-03-02

PRIOR APPLICATION NUMBER: 60/039,762

PRIOR FILING DATE: 1997-03-03

NUMBER OF SEQ ID NOS: 71

SOFTWARE: PatentIn Ver. 2.0

SEQ ID NO: 44

LENGTH: 822

TYPE: DNA

ORGANISM: Unknown

FEATURE:

; OTHER INFORMATION: Description of Unknown Organism: unknown

US-10-139-089-44

Query Match 100 %; Score 822; DB 16; Length 822;

Best Local Similarity 100 %; Pred. No. 1.e-216;

Matches 822; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 CGATGCTGAACTGTACTTAGACTAACTTGAGCAATAACACAGTAGTGTGAG 60

Db 1 GCATGCTGAACTGTACTTAGACTAACTTGAGCAATAACACAGTAGTGTGAG 60

QY 61 GATTGTTGCCTGGTACATCACACTCTGTCAGTCAGCTCTCTTGTCTGG 120

Db 121 AAATTGCTGTTCTGTTGTTCTCCTTCACTGCTATCTATTTCTAACCTCA 180

Db 121 AAATTGCTGTTCTGTTGTTCTCCTTCACTGCTATCTATTTCTAACCTCA 180

QY 181 CATGCCATACAATCTGCTGCAAGCTATGATTCCTCCAAATATCTATCTATTTCTAACCTCA 180

Db 181 CATGCCATACAATCTGCTGCAAGCTATGATTCCTCCAAATATCTATCTATTTCTAACCTCA 180

QY 241 CTTGTCACAGATAAAAGTAGTATCAATGCACTAACGCTTCACTTGAGGSC 300

Db 241 CTTGTCACAGATAAAAGTAGTATCAATGCACTAACGCTTCACTTGAGGSC 300

QY 301 TTAAGCAGTTCAACATACAAACCGGGAGTTGCCCTGAATGTTCTCAAATGTT 360

Db 301 TTAAGCAGTTCAACATACAAACCGGGAGTTGCCCTGAATGTTCTCAAATGTT 360

QY 361 CCTGTAACATAGGGCTCTGTCTTAACATAGTAGTATCAATGCACTGGCTCA 420

Db 361 CCTGTAACATAGGGCTCTGTCTTAACATAGTAGTATCAATGCACTGGCTCA 420

QY 421 TCCCACTATGGGGATGAGATGAAAGGGAGCCTGAATTAATTTACATAGTCAA 480

Db 421 TCCCACTATGGGGATGAGATGAAAGGGAGCCTGAATTAATTTACATAGTCAA 480

QY 481 TAGGATAGCCAGGCTTGGTAATCTGCCTATCTAATATTC 540

Db 481 TAGGATAGCCAGGCTTGGTAATCTGCCTATCTAATATTC 540

QY 541 CAAACTGAACTGACTGACTGTTAGTACTAGTCTTCACTAAATTC 540

Db 541 CAAACTGAACTGACTGACTGTTAGTACTAGTCTTCACTAAATTC 540

QY 601 CTTATCCAGGCCACTTATGAGCTGTGCTCTGAACTAAATACCGTAT 660

Db 601 CTTATCCAGGCCACTTATGAGCTGTGCTCTGAACTAAATACCGTAT 660

QY 661 GCTGTTAATTATGCAATGTCATTTCAACCTAGAAATACCTTAAGTACAGA 720

Db 661 GCTGTTAATTATGCAATGTCATTTCAACCTAGAAATACCTTAAGTACAGA 720

QY 721 TATACACAAAGGTTACTGTTACGGATGCTGTGAAAGAGTAAAGATTTC 780

Db 721 TATACACAAAGGTTACTGTTACGGATGCTGTGAAAGAGTAAAGATTTC 780

QY 781 AGCATGATTCCATATGTGCTTCCACCTACTGCCAATACCA 822

Db 781 AGCATGATTCCATATGTGCTTCCACCTACTGCCAATACCA 822

RESULT 9
US-10-822-873-6

; Sequence 6, Application US/10/822-873

; Publication No. US2004024185A1

; GENERAL INFORMATION:

; APPLICANT: Henderson, D.R.

; APPLICANT: Schuur, E.R.

TITLE OF INVENTION: TISSUE SPECIFIC ADENOVIRAL VECTORS

FILE REFERENCE: CELL-004CON2

CURRENT APPLICATION NUMBER: US/10/822,873

PRIOR APPLICATION NUMBER: 09/732,169

PRIOR FILING DATE: 2000-12-06

PRIOR APPLICATION NUMBER: 09/151,376

PRIOR FILING DATE: 1998-09-10

PRIOR APPLICATION NUMBER: 08/669,753

PRIOR FILING DATE: 1996-01-26

PRIOR APPLICATION NUMBER: 08/495,034

PRIOR FILING DATE: 1995-06-27

NUMBER OF SEQ ID NOS: 71

SOFTWARE: PatentIn Ver. 2.0

APPLICATION: Schuur, E.R.
 TITLE OF INVENTION: TISSUE SPECIFIC ADENOVIRAL VECTORS
 FILE REFERENCE: CELL-004CON2
 CURRENT APPLICATION NUMBER: US/10/822,873
 CURRENT FILING DATE: 2004-04-13
 PRIOR APPLICATION NUMBER: 09/732,169
 PRIOR FILING DATE: 2000-12-06
 PRIOR APPLICATION NUMBER: 09/151,376
 PRIOR FILING DATE: 1998-09-10
 PRIOR APPLICATION NUMBER: 08/669,753
 PRIOR FILING DATE: 1996-06-26
 PRIOR APPLICATION NUMBER: 08/495,034
 PRIOR FILING DATE: 1995-06-27
 NUMBER OF SEQ ID NOS: 71
 SOFTWARE: PatentIn Ver. 2.0
 SEQ ID NO: 44
 LENGTH: 822
 TYPE: DNA
 ORGANISM: Unknown
 FEATURE:
 OTHER INFORMATION: Description of Unknown Organism: unknown
 US-10-822-873-44

Query Match 100.0%; Score 822; DB 20; Length 822;
 Best Local Similarity 100.0%; Pred. No. 1.1e-216; Mismatches 0; Indels 0; Gaps 0;
 Matches 822; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 GCATGGCTGTAACCTGTGACTTGTGACTTAGGACTAACTTGAGCAATAACACATAGATGAG 60
 Db 1 GCATGGCTGTAACCTGTGACTTGTGACTTAGGACTAACTTGAGCAATAACACATAGATGAG 60
 QY 61 GATTTGTTGCTGTGAGCATACAAGCTCTGCTGAAAGCTTGTGACTTAGGACTAACTTGAGCAATAACACATAGATGAG 60
 Db 61 GATTTGTTGCTGTGACTTGTGACTTAGGACTAACTTGAGCAATAACACATAGATGAG 60
 QY 121 AAAATTGCTGTCATGTTCTTCATGCTTCACTGCTATCTATTTCTAACACTCA 180
 Db 121 AAAATTGCTGTCATGTTCTTCATGCTTCACTGCTATCTATTTCTAACACTCA 180
 QY 181 CATGGCTACATAACTGTCTGCAAGCTTGTGACTTAGGACTAACTTGAGCAATAACACATAGATGAG 240
 Db 181 CATGGCTACATAACTGTCTGCAAGCTTGTGACTTAGGACTAACTTGAGCAATAACACATAGATGAG 240
 QY 241 CTTGCTCAGAAGATAAAAGTACTATCAATGCAATCACGCTCCACTTGAGGCC 300
 Db 241 CTTGCTCAGAAGATAAAAGTACTATCAATGCAATCACGCTCCACTTGAGGCC 300
 QY 301 TTAAAGACCTTCACATACAAACCGGGGAGTTGCCTGAATTTCTTAAMAGTG 360
 Db 301 TTAAAGACCTTCACATACAAACGGGGAGTTGCCTGAATTTCTTAAMAGTG 360
 QY 361 CCTGTAGCACATAGGTCTCTGTTCTTAAATCTAACTTTAGGCCAGTGCTCA 420
 Db 361 CCTGTAGCACATAGGTCTCTGTTCTTAAATCTAACTTTAGGCCAGTGCTCA 420
 QY 421 TCCACCTATGGGAGATGAGCTGAAAGGGCCCTGATTAATTTACAGTCAA 480
 Db 421 TCCACCTATGGGAGATGAGCTGAAAGGGCCCTGATTAATTTACAGTCAA 480
 QY 481 TAGGATAGGCCAGGACTGTTGGTAACTGGTCACTTATTAACATAATATAC 540
 Db 481 TAGGATAGGCCAGGACTGTTGGTAACTGGTCACTTATTAACATAATATAC 540
 QY 541 CAAACTGACATGACTGTTAGTTACTAAGCTTATCTCATACATACCTCG 600
 Db 541 CAAACTGACATGACTGTTAGTTACTAAGCTTATCTCATACATACCTCG 600
 QY 601 CTTTATCCAGGCCCTATGAGCTGTCCTGAACTAAATACAAATACCGCTAT 660
 Db 601 CTTTATCCAGGCCCTATGAGCTGTCCTGAACTAAATACAAATACCGCTAT 660
 QY 661 GCTGTTAATATGGCAATGTCCTTCAACCTAACGAAATACATAAGTACAGA 720
 Db 661 GCTGTTAATATGGCAATGTCCTTCAACCTAACGAAATACATAAGTACAGA 720
 QY 721 TATACACAAAGGTTACTAGTACAGGCTGCTGAAAGGATATAAGGATTG 780
 Db 721 TATACACAAAGGTTACTAGTACAGGCTGCTGAAAGGATATAAGGATTG 780
 QY 781 AGCATGTTCCATATGTCCTCACCACTGCCATACA 822
 Db 781 AGCATGTTCCATATGTCCTCACCACTGCCATACA 822

RESULT 10
 US-10-822-873-44
 Sequence 44, Application US/10822873
 Publication No. US20040241857A1
 GENERAL INFORMATION:
 APPLICANT: Henderson, D.R.

RESULT 11
US-10-691-045-8
; Sequence 8, Application US/10691045
; Publication No. US20040146489A1
; GENERAL INFORMATION:
; APPLICANT: Yu, De-Chao
; APPLICANT: Li, Yuanbao
; APPLICANT: Henderson, Daniel R.
; TITLE OF INVENTION: CELL-SPECIFIC ADENOVIRUS VECTORS
; TITLE OF INVENTION: COMPRISING AN INTERNAL RIBOSOME ENTRY SITE
; PILE REFERENCE: 34802201700
; CURRENT APPLICATION NUMBER: US/10/691,045
; CURRENT FILING DATE: 2003-10-21
; PRIOR APPLICATION NUMBER: US/07/814,351
; PRIOR FILING DATE: 2001-03-21
; NUMBER OF SEQ ID NOS: 35
; SOFTWARE: fastSEQ for Windows Version 4.0
; SEQ ID NO 8
; LENGTH: 822
; TYPE: DNA
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: APP-TRE

Query Match 100.0% Score 822; DB 22; Length 822;
Best Local Similarity 100.0%; Pred. No. 1.1e-216; Mismatches 0; Indels 0; Gaps 0;
Matches 822; Conservative 0;

QY 1 GCATTTGCTGTGAACTCTGTACTTGTAGCAACTAACACAGCTAGTAGGAG 60
Db 1 GCATTTGCTGTGAACTCTGTACTTGTAGCAACTAACACAGCTAGTAGGAG 60
QY 61 GATTGTTGCTGTAGCATGAACTCTGGTCAAGCTCCTTATGCTGCTGG 120
Db 61 GATTGTTGCTGTAGCATGAACTCTGGTCAAGCTCCTTATGCTGCTGG 120
QY 121 AATTTGCTGTCTCATGGTTCTTCATCTTCTACTGCTATCTTCTACCACTCA 180
Db 121 AATTTGCTGTCTCATGGTTCTTCATCTTCTACTGCTATCTTCTACCACTCA 180
QY 181 CATGGCTACATAACTGTCGAACGTTAGTATCCAAATATCTCTAGCTCAAT 240
Db 181 CATGGCTACATAACTGTCGAACGTTAGTATCCAAATATCTCTAGCTCAAT 240
QY 241 CTGGTCCAGAGATAAAAGTAGTTCAATGCATCAGTCAGCTCCAGTGAGGGC 300
Db 241 CTGGTCCAGAGATAAAAGTAGTTCAATGCATCAGTCAGCTCCAGTGAGGGC 300
QY 301 TAAACAGCTTCAACATACAAACCGGGAGCTTGCCTGCGATGTTCCAAATGTT 360
Db 301 TAAACAGCTTCAACATACAAACCGGGAGCTTGCCTGCGATGTTCCAAATGTT 360
QY 361 CCTGTTGCACTTAGGTTCTCTGCTTAATCTTGTAGCTTGTGCTCA 420
Db 361 CCTGTTGCACTTAGGTTCTCTGCTTAATCTTGTAGCTTGTGCTCA 420
QY 421 TCCCACTATGGGAGATGAGGTGAAAGGGAGCTGTTAAATACAACTAAGTCAA 480
Db 421 TCCCACTATGGGAGATGAGGTGAAAGGGAGCTGTTAAATACAACTAAGTCAA 480

RESULT 12
US-09-828-455-12
; Sequence 12, Application US/09828455
; Publication No. US20030106621A1
; GENERAL INFORMATION:
; APPLICANT: TAMOKI, TAKI
; TITLE OF INVENTION: IMPROVED METHOD OF EXPRESSING GENES IN
; NUMBER OF SEQUENCES: 46
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: BURNS, DOANE, SWECKER & MATHIS, L.L.P.
; STREET: 699 PRINCE STREET
; CITY: ALEXANDRIA
; STATE: VA
; COUNTRY: USA
; ZIP: 22313-1404
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPILER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/09/828,455
; FILING DATE: 09-APR-2001
; CLASSIFICATION: 435
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US/08/645,215
; FILING DATE: 13-MAY-1996
; APPLICATION NUMBER: US/08/148,058
; FILING DATE: 04-NOV-1993
; ATTORNEY/AGENT INFORMATION:
; NAME: MOOI, LESLIE A.
; REGISTRATION NUMBER: 37,047
; REFERENCE/DOCKET NUMBER: 028722-135
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 415-854-7400
; TELEFAX: 415-854-8275
; INFORMATION FOR SEQ ID NO: 12:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 2240 base pairs
; TYPE: nucleic acid
; STRANDBNESS: double
; TOPOLOGY: linear
; SEQUENCE DESCRIPTION: SEQ ID NO: 12:
; US-09-828-455-12

Query Match 75.3%; Score 619; DB 10; Length 2240;
 Best Local Similarity 100.0%; Pred. No. 2.3e-160; Mismatches 0; Indels 0; Gaps 0; Matches 619; Conservative 0; MisMatches 0; Indels 0; Gaps 0;

Y 1 GCATTCGTGTAACCTGTACTAGGACTAACTTGAGCAATACACATAGATTGAG 60
 Db 1242 GATTTGCGTGAACCTGTACTAGGACTAACTTGAGCAATACACATAGATTGAG 1301

Y 61 GATTTGCGTGAACCTGTACTAGGACTAACTTGAGCAATACACATAGATTGAG 60
 Db 1302 GATTTGCGTGAACCTGTACTAGGACTAACTTGAGCAATACACATAGATTGAG 1361

Y 121 AAATTGCTGTTCTATGGTTCTTCACTGCTATCTATTTCTCAACCCTA 180
 Db 1362 AAATTGCTGTTCTATGGTTCTATGGCTAAGCTTAATGATTCCAAATATCTATCTAGCTCAAT 240

Y 1422 CATGGCTCACATACTGCTGCAAGCTTATGATCCAAATATCTATCTAGCTCAAT 1481

Y 241 CTGTTCCAGAGATAAAAGTAGTATCAATGCCAACGTTCCACTGGAGGC 300
 Db 1482 CTGTTCCAGAGATAAAAGTAGTATCAATGCCAACGTTCCACTGGAGGC 1541

Y 301 TTAAGAGCTTCACATCACACACACACGGGGAGTTGGCTCGAAGTTTCCTAAATGTT 360

Y 1542 TTAAGAGCTTCACATCACACACACGGGGAGTTGGCTCGAAGTTTCCTAAATGTT 1601

Y 361 CCTGTCACATAGGGCTCTGTTCTTCAAATCTTAATCTTACTTTAGCCAGTGCTCA 420
 Db 1602 CCTGTCACATAGGGCTCTGTTCTTCAAATCTTAATCTTACTTTAGCCAGTGCTCA 1661

Y 421 TCCACCATGCGGAGTGAGCTGAAAGGGGCCGATTAACTTACACTAGTC 480
 Db 1662 TCCACCATGCGGAGTGAGCTGAAAGGGGCCGATTAACTTACACTAGTC 1721

Y 481 TAGGATAGAGCCAGGCTGTTGGTAACTGGTCACTTATCTTAACTAATATC 540
 Db 1722 TAGGATAGAGCCAGGCTGTTGGTAACTGGTCACTTATCTTAACTAATATC 1781

Y 541 CAAACTGAACTGACTTAGTACTAGTCTGTTCACTTACCACTCTAG 600
 Db 1782 CAAACTGAACTGACTTAGTACTAGTCTGTTCACTTACCACTCTAG 1841

Y 601 CTTTATCCAGGCCACTAT 619
 Db 1842 CTTTATCCAGGCCACTAT 1860

RESULT 13
 S-09-898-893-2
 Sequence 2, Application US/09898883
 Patent No. US20020164793A1
 GENERAL INFORMATION:
 APPLICANT: Little, Andrew
 Lamparski, Henry
 Schuur, Eric
 Henderson, Daniel
 TITLE OF INVENTION: ABDONVIRUS VECTORS SPECIFIC FOR CELLS EXPRESSING ALPHA-FETOPROTEIN AND METHODS OF USE THEREOF
 NUMBER OF SEQUENCES: 23
 CORRESPONDENCE ADDRESS:
 ADDRESSEE: MORRISON & FOERSTER
 STREET: 755 PAGE MILL ROAD
 CITY: PALO ALTO
 STATE: CA
 COUNTRY: USA
 ZIP: 94304-1018
 COMPUTER READABLE FORM:
 MEDIUM TYPE: floppy disk
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: PatentIn Release #1.0, Version #1.30

CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/09/898, 883
 FILING DATE: 02-nul-2001
 CLASSIFICATION: <Unknown>
 PRIORITY APPLICATION DATA:
 APPLICATION NUMBER: US/09/033, 428
 FILING DATE: <Unknown>
 ATTORNEY/AGENT INFORMATION:
 NAME: POLIZZI, CATHERINE M.
 REGISTRATION NUMBER: 40,130
 REFERENCE/DOCKET NUMBER: 34802-3-0004.00
 TELECOMMUNICATION INFORMATION:
 TELEPHONE: (415) 813-5600
 TELEFAX: (415) 494-0792
 TELEX: 706141 MRSNFOERS SFO
 SEQUENCE CHARACTERISTICS:
 LENGTH: 5224 base pairs
 TYPE: nucleic acid
 STRANDEDNESS: single
 TOPOLOGY: linear
 SEQUENCE DESCRIPTION: SEQ ID NO: 2:
 US-09-898-883-2

Query Match 75.3%; Score 619; DB 9; Length 5224;
 Best Local Similarity 100.0%; Pred. No. 4.3e-160; Mismatches 0; Indels 0; Gaps 0; Matches 619; Conservative 0; MisMatches 0; Indels 0; Gaps 0;

Qy 1 GATTTGCGTGTCTTCACTGCTTCTCTTACTGCTATCTTCTTCAACCCTA 180
 Db 1242 GATTTGCGTGTCTTCACTGCTTCTCTTACTGCTATCTTCTTCAACCCTA 1301

Qy 61 GATTTGCGTGTCTTCACTGCTTCTCTTACTGCTATCTTCTTCAACCCTA 180
 Db 1302 GATTTGCGTGTCTTCACTGCTTCTCTTACTGCTATCTTCTTCAACCCTA 1361

Qy 121 AAATTGCTGTTCTATGGTTCTTCACTGCTATCTTACTGCTATCTTCTTCAACCCTA 180
 Db 1362 AAATTGCTGTTCTATGGTTCTTCACTGCTATCTTACTGCTATCTTCTTCAACCCTA 1421

Qy 181 CATGGCTCACATACTGCTTCACTTACCACTCTAGCTTATCTTCACTTACCACT 240
 Db 1422 CATGGCTCACATACTGCTTCACTTACCACTCTAGCTTATCTTCACTTACCACT 1481

Qy 241 CTGTTCCAGAGATAAAAGTAGTATCAATGCCAACGTTCCACTGGAGGC 300
 Db 1482 CTGTTCCAGAGATAAAAGTAGTATCAATGCCAACGTTCCACTGGAGGC 1541

Qy 301 TTAAGAGCTTCACATCACACACACGGGGAGTTGGCTGGAAAGTTCTTAAATGTT 360
 Db 1542 TTAAGAGCTTCACATCACACACGGGGAGTTGGCTGGAAAGTTCTTAAATGTT 1601

Qy 361 CCTGTCACATAGGGCTCTGTTCTTCAAATCTTAATCTTAACTTCTTAACTAATATC 420
 Db 1602 CCTGTCACATAGGGCTCTGTTCTTCAAATCTTAATCTTAACTTCTTAACTAATATC 1661

Qy 421 TCCACCATGCGGAGTGAGCTGAAAGGGGCCGATTAACTTACACTAGTC 480
 Db 1662 TCCACCATGCGGAGTGAGCTGAAAGGGGCCGATTAACTTACACTAGTC 1721

Qy 481 TAGGATAGAGCCAGGCTGTTGGTAACTGGTCACTTATCTTAACTAATATC 540
 Db 1722 TAGGATAGAGCCAGGCTGTTGGTAACTGGTCACTTATCTTAACTAATATC 1781

Qy 541 CAAACTGAACTGACTTAGTACTAGTCTGTTCACTTACCACTCTAG 600
 Db 1782 CAAACTGAACTGACTTAGTACTAGTCTGTTCACTTACCACTCTAG 1841

Qy 601 CTTTATCCAGGCCACTAT 619
 Db 1842 CTTTATCCAGGCCACTAT 1860

RESULT 14
US-09-151-376-5
Sequence 5, Application US/09151376
Publication No. US20030044383A1
; GENERAL INFORMATION:
; APPLICANT: Henderson, D.R.
; APPLICANT: Schuur, E.R.
; TITLE OF INVENTION: TISSUE SPECIFIC VIRAL VECTORS
; FILE REFERENCE: 348022000221
; CURRENT APPLICATION NUMBER: US/09/151,376
; CURRENT FILING DATE: 1998-09-10
; EARLIER APPLICATION NUMBER: 08/669,753
; EARLIER FILING DATE: 1996-06-26
; EARLIER APPLICATION NUMBER: 08/495,034
; EARLIER FILING DATE: 1995-06-27
; NUMBER OF SEQ ID NOS: 71
; SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 5
LENGTH: 5224
TYPE: DNA
ORGANISM: Homo sapiens
; US-09-151-376-5
Query Match 75.3%; Score 619; DB 10; Length 5224;
Best Local Similarity 100.0%; Pred. No. 4.3e-160; Matches 619; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
QY 1 GCATGGTGTGAACTCTGTACTTGTAGGACTAACTTGAGCATACACATAGATTGAG 60
Db 1242 GCATGGTGTGAACTCTGTACTTGTAGGACTAACTTGAGCATACACATAGATTGAG 1301
QY 61 GATTTGTTGCTTGTAGCATACAACTCTGTCAAGCTCCTTATGTTGCTTGG 120
Db 1302 GATTTGTTGCTTGTAGCATACAACTCTGTCAAGCTCCTTATGTTGCTTGG 1361
QY 121 AAATTGTCGTTCTCATGTTCTTCACTGTTACTGCTATCTTGTCAACACTCA 180
Db 1362 AAATTGTCGTTCTCATGTTCTTCACTGTTACTGCTATCTTGTCAACACTCA 1421
QY 181 CATGGTACATAACTGTCGAAGTTATGTTCCAAATATCTCTAGCTCAAT 240
Db 1422 CATGGTACATAACTGTCGAAGTTATGTTCCAAATATCTCTAGCTCAAT 1481
QY 241 CTTGTCCAGAGATAAAAGTAGTATCAGTCATCACAGTGTCTCACTGGAGGC 300
Db 1482 CTTGTCCAGAGATAAAAGTAGTATCAGTCATCACAGTGTCTCACTGGAGGC 1541
QY 301 TTAAGAGCGTTCAACATACAAACCGGGAGTTTCCTGAAATGTTCCAAATGTT 360
Db 1542 TTAAGAGCGTTCAACATACAAACCGGGAGTTTCCTGAAATGTTCCAAATGTT 1601
QY 361 CCTGTAGACATAGGGCTCTGTCTTAACATTAATCTTACTTTAGGCCAGGTCA 420
Db 1602 CCTGTAGACATAGGGCTCTGTCTTAACATTAATCTTACTTTAGGCCAGGTCA 1661
QY 421 TCCACCTATGGGAGATGAGTGAAGGGAGCTGATTATACTCTAAGTCAA 480
Db 1662 TCCACCTATGGGAGATGAGTGAAGGGAGCTGATTATACTCTAAGTCAA 1721
QY 481 TAGGCATAGACCCAGGACTGTTGGTAACCTGGTACTTTACCTTAACCTAATATC 540
Db 1722 TAGGCATAGACCCAGGACTGTTGGTAACCTGGTACTTTACCTTAACCTAATATC 1781
QY 541 CAAACTGAACTGATGTTACTGACTTACCTTAACCTAATATC 600
Db 1782 CAAACTGAACTGATGTTACTGACTTACCTTAACCTAATATC 1841
QY 601 CTTATCCAGCCACTTAT 619
Db 1842 CTTATCCAGCCACTTAT 1860
; US-09-151-376-5
Query Match 75.3%; Score 619; DB 10; Length 5224;
Best Local Similarity 100.0%; Pred. No. 4.3e-160; Matches 619; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
QY 1 GCATGGTGTGAACTCTGTACTTGTAGGACTAACTTGAGCATACACATAGATTGAG 60
Db 1242 GCATGGTGTGAACTCTGTACTTGTAGGACTAACTTGAGCATACACATAGATTGAG 1301
QY 61 GATTTGTTGCTTGTAGCATACAACTCTGTCAAGCTCCTTATGTTGCTTGG 120
Db 1302 GATTTGTTGCTTGTAGCATACAACTCTGTCAAGCTCCTTATGTTGCTTGG 1361
QY 121 AAATTGTCGTTCTCATGTTCTTCACTGTTACTGCTATCTTGTCAACACTCA 180
Db 1362 AAATTGTCGTTCTCATGTTCTTCACTGTTACTGCTATCTTGTCAACACTCA 1421
QY 181 CATGGTACATAACTGTCGAAGTTATGTTCCAAATATCTCTAGCTCAAT 240
Db 1422 CATGGTACATAACTGTCGAAGTTATGTTCCAAATATCTCTAGCTCAAT 1481
QY 241 CTTGTCCAGAGATAAAAGTAGTATCAGTCATCACAGTGTCTCACTGGAGGC 300
Db 1482 CTTGTCCAGAGATAAAAGTAGTATCAGTCATCACAGTGTCTCACTGGAGGC 1541
QY 301 TTAAGAGCGTTCAACATACAAACCGGGAGTTTCCTGAAATGTTCCAAATGTT 360
Db 1542 TTAAGAGCGTTCAACATACAAACCGGGAGTTTCCTGAAATGTTCCAAATGTT 1601
QY 361 CCTGTAGACATAGGGCTCTGTCTTAACATTAATCTTACTTTAGGCCAGGTCA 420
Db 1602 CCTGTAGACATAGGGCTCTGTCTTAACATTAATCTTACTTTAGGCCAGGTCA 1661
QY 421 TCCACCTATGGGAGATGAGTGAAGGGAGCTGATTATACTCTAAGTCAA 480
Db 1662 TCCACCTATGGGAGATGAGTGAAGGGAGCTGATTATACTCTAAGTCAA 1721
QY 481 TAGGCATAGACCCAGGACTGTTGGTAACCTGGTACTTTACCTTAACCTAATATC 540
Db 1722 TAGGCATAGACCCAGGACTGTTGGTAACCTGGTACTTTACCTTAACCTAATATC 1781
QY 541 CAAACTGAACTGATGTTACTGACTTACCTTAACCTAATATC 600
Db 1782 CAAACTGAACTGATGTTACTGACTTACCTTAACCTAATATC 1841
QY 601 CTTATCCAGCCACTTAT 619
Db 1842 CTTATCCAGCCACTTAT 1860

Mon Aug 8 10:19:28 2005

us-09-509-591-1.rnpb

Page 11

Search completed: August 7, 2005, 08:15:56
Job time : 495.167 secs

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OM nucleic - nucleic search, using SW model

Run on: August 6, 2005, 16:35:01 ; Search time 127.256 Seconds
(without alignments)
10569.369 Million cell updates/sec

Title: US-09-509-591-1
Perfect score: 822
Sequence: 1 GCATGCTGAACTCTGTAA.....TTCCACCACTGCCATAACA 822

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 1202784 seqs, 818138359 residues

Total number of hits satisfying chosen parameters: 2405568

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Issued Patents NA:
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2: /cgn2_6/pcdodata/1/ina/5B_COMB.seq:
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

ALIGNMENTS

RESULT 1
US-09-033-428-1
; Sequence 1, Application US/09033428
; Patent No. 6254862

; GENERAL INFORMATION:
; APPLICANT: Little, Andrew
; APPLICANT: Lamparski, Henry
; APPLICANT: Schuur, Eric
; APPLICANT: Henderson, Daniel
; TITLE OF INVENTION: ADENOVIRUS VECTORS SPECIFIC FOR CELLS
; TITLE OF INVENTION: EXPRESSING ALPHA-FETOPROTEIN AND METHODS OF USE THEREOF
; NUMBER OF SEQUENCES: 23
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: MORRISON & FOERSTER
; STREET: 755 PAGE MILL ROAD
; CITY: PALO ALTO
; STATE: CA
; COUNTRY: USA
; ZIP: 94304-1018

; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/09/033,428
; FILING DATE:
; CLASSIFICATION:
; ATTORNEY/AGENT INFORMATION:
; NAME: POLIZZI, CATHERINE M.
; REGISTRATION NUMBER: 40,130
; TELEFAX: (415) 813-5600
; TELEX: 706141 MRSNFO SFO
; INFORMATION FOR SEQ ID NO: 1:
; REFERENCE/DOCKET NUMBER: 34802-30004.00
; SEQUENCE CHARACTERISTICS:
; LENGTH: 822 base Pairs
; TELEPHONE: (415) 813-5600
; CLASSIFICATION:
; TYPE: nucleic acid
; STANDINGNESS: Single
; TOPOLOGY: Linear

US-09-033-428-1

SUMMARIES

| Result No. | Score | Query | Match | Length | DB | ID | Description |
|------------|-------|-------|--------|--------|---------------------|---------------------|---------------------|
| 1 | 822 | 100.0 | 822 | 3 | US-09-033-428-1 | Sequence 1, Appli | Sequence 1, Appli |
| 2 | 822 | 100.0 | 822 | 3 | US-09-033-556-6 | Sequence 6, Appli | Sequence 6, Appli |
| 3 | 822 | 100.0 | 822 | 4 | US-09-474-699-14 | Sequence 14, Appli | Sequence 14, Appli |
| 4 | 822 | 100.0 | 822 | 4 | US-09-898-883-1 | Sequence 1, Appli | Sequence 1, Appli |
| 5 | 822 | 100.0 | 822 | 4 | US-09-151-376-6 | Sequence 6, Appli | Sequence 6, Appli |
| 6 | 822 | 100.0 | 822 | 4 | US-09-151-376-44 | Sequence 44, Appli | Sequence 44, Appli |
| 7 | 822 | 100.0 | 822 | 4 | US-09-814-351-8 | Sequence 8, Appli | Sequence 8, Appli |
| 8 | 619 | 75.3 | 2240 | 1 | US-08-148-05A-12 | Sequence 12, Appli | Sequence 12, Appli |
| 9 | 619 | 75.3 | 2240 | 1 | US-08-478-042-12 | Sequence 12, Appli | Sequence 12, Appli |
| 10 | 619 | 75.3 | 2240 | 1 | US-08-645-215-12 | Sequence 12, Appli | Sequence 12, Appli |
| 11 | 619 | 75.3 | 2240 | 2 | US-08-166-604-12 | Sequence 12, Appli | Sequence 12, Appli |
| 12 | 619 | 75.3 | 5224 | 3 | US-09-033-428-2 | Sequence 2, Appli | Sequence 2, Appli |
| 13 | 619 | 75.3 | 5224 | 3 | US-09-556-6 | Sequence 5, Appli | Sequence 5, Appli |
| 14 | 619 | 75.3 | 5224 | 4 | US-09-774-699-15 | Sequence 15, Appli | Sequence 15, Appli |
| 15 | 619 | 75.3 | 5224 | 4 | US-09-988-803-2 | Sequence 2, Appli | Sequence 2, Appli |
| 16 | 619 | 75.3 | 5224 | 4 | US-09-151-376-5 | Sequence 5, Appli | Sequence 5, Appli |
| 17 | 619 | 75.3 | 5224 | 4 | US-09-151-376-45 | Sequence 45, Appli | Sequence 45, Appli |
| 18 | 619 | 75.3 | 5224 | 4 | US-09-914-351-16 | Sequence 16, Appli | Sequence 16, Appli |
| 19 | 203.6 | 24.8 | 23497 | 4 | US-09-949-016-17072 | Sequence 17072, A | Sequence 17072, A |
| 20 | 73 | 8.9 | 228 | 3 | US-08-181-96A-3 | Sequence 3, Appli | Sequence 3, Appli |
| 21 | 73 | 8.9 | 228 | 3 | US-08-154-712B-3 | Sequence 3, Appli | Sequence 3, Appli |
| 22 | 73 | 8.9 | 228 | 4 | US-09-947-95A-3 | Sequence 17216, A | Sequence 17216, A |
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| 24 | 50 | 6.1 | 50 | 1 | US-08-171-389-20 | Sequence 352, Appli | Sequence 352, Appli |
| 25 | 50 | 6.1 | 50 | 1 | US-08-123-349-352 | Sequence 20, Appli | Sequence 20, Appli |
| 26 | 50 | 6.1 | 50 | 1 | US-08-123-916-20 | Sequence 352, Appli | Sequence 352, Appli |
| 27 | 50 | 6.1 | 50 | 1 | US-08-123-916-352 | Sequence 14619, A | Sequence 14619, A |

Query Match Score 822; DB 3; Length 822;
Best Local Similarity 100.0%; Pred. No. 1.5-223;
Matches 822; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
Qy 1 GCATGCTGAACTCTGTACTTAGGACTAACITGAGGATACACATAGTTGAG 60

Page 2

Db 601 CTTTATCAGGCCACATTAGAGCTCTGGTCACCTTAACAAATAACCGCTAT 660 Qy 661 GCTGTTAATTATGGCAATGTCGCAATTCAACCTAAACTAAGTAACGGA 720 Db 661 GCTGTTAATTATGGCAATGTCGCAATTCAACCTAAACTAAGTAACGGA 720 Qy 721 TATACCAAAAGGTACTAGTTAACGGCATGGCTGAAGAGTTAAAGAATTTC 780 Db 721 TATACCAAAAGGTACTAGTTAACGGCATGGCTGAAGAGTTAAAGAATTTC 780 Qy 781 AGCATGATTTCATATTTGTGTCACACTGCCAAATAACA 822 Db 781 AGCATGATTTCATATTTGTGTCACACTGCCAAATAACA 822 Qy 781 AGCATGATTTCATATTTGTGTCACACTGCCAAATAACA 822

RESULT 3
US-09-474-699-14
; Sequence 1, Application US/09474699
; Patent No. 6495130
; GENERAL INFORMATION:
; APPLICANT: Henderson, Daniel R.
; TITLE OF INVENTION: TARGET CELL-SPECIFIC ADENOVIRAL VECTORS
; TITLE OF INVENTION: CONTAINING E3 AND METHODS OF USE THEREOF
; FILE REFERENCE: 348022001300
; CURRENT APPLICATION NUMBER: US/09/474-699
; CURRENT FILING DATE: 1999-12-29
; PRIOR APPLICATION NUMBER: 60/114,262
; PRIOR FILING DATE: 1998-12-30
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO: 14
; LENGTH: 822
; TYPE: DNA
; ORGANISM: Homo Sapien
US-09-474-699-14

Query Match 100.0%; Score 822; DB 4; Length 822;
Best Local Similarity 100.0%; Pred. No. 1.5e-223; Matches 822; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 GCATGGTGTGAACCTCTGACTTCTAGGCTAAACTTGAACATAACAGATTGAG 60 Db 1 GCATGGTGTGAACCTCTGACTTCTAGGCTAAACTTGAACATAACAGATTGAG 60

Qy 1 AAAATTGCTGTCTCTGGTTCTCTTGACTGTTCTTCACTGTTCTACCTCA 180 Db 121 AAAATTGCTGTCTCTGGTTCTCTTGACTGTTCTTCACTGTTCTACCTCA 180

Qy 181 CATGGCTAACATAACTCTGCAAGCTTATGATTCCAAATPATCTCATCTGCTCAAT 240 Db 181 CATGGCTAACATAACTCTGCAAGCTTATGATTCCAAATPATCTCATCTGCTCAAT 240

Qy 241 CTTGTTCAGAGATAAAAGTAGTATTCAATGCCATCAACGCTCTCCTTGAGGGC 300 Db 241 CTTGTTCAGAGATAAAAGTAGTATTCAATGCCATCAACGCTCTCCTTGAGGGC 300

Qy 301 TTAAAGAGCTTCACATACAAACCGGGAGTTTGCCTGAAATGTTCTPAAATGT 360 Db 301 TTAAAGAGCTTCACATACAAACCGGGAGTTTGCCTGAAATGTTCTPAAATGT 360

Qy 361 CCTGTACACATAGGGCTCTTGTCTTAAATCTAAATFACTTTAGCCAGTNGCTCA 420 Db 361 CCTGTACACATAGGGCTCTTGTCTTAAATCTAAATFACTTTAGCCAGTNGCTCA 420

Qy 421 TCCCACCTATGGGAGATGAGGTGAAAGGGAGCTGATTAATTAACTAGTCAA 480 Db 421 TCCCACCTATGGGAGATGAGGTGAAAGGGAGCTGATTAATTAACTAGTCAA 480

481 TAGGCATAGGCCAGGACTGTTGGTAAACTCTGTCACTTTATCTTAAACTAATATTC 540
481 TAGGCATAGGCCAGGACTGTTGGTAAACTCTGTCACTTTATCTTAAACTAATATTC 540
541 CAAACTGAAACATGTTACTCTAGTTACTAAGTCCTTGCCTTACATCACCCTCAG 600
541 CAAACTGAAACATGTTACTCTAGTTACTAAGTCCTTGCCTTACATCACCCTCAG 600
601 CTTTATCAGGCCACCTATGAGCTCTGTCCTGACATAAAATCAAATAACCGCTAT 660
601 CTTTATCAGGCCACCTATGAGCTCTGTCCTGACATAAAATCAAATAACCGCTAT 660
661 GCTGTTAATTATTGGCAATGTCACCTTCAACCTAAGAACTAACAGA 720
661 GCTGTTAATTATTGGCAATGTCACCTTCAACCTAAGAACTAACAGA 720
Qy 721 TATACCAAAAGGTACTAGTTAACGGCATTTGCTGAAAGAGCTTAAAGAATTTC 780
Db 721 TATACCAAAAGGTACTAGTTAACGGCATTTGCTGAAAGAGCTTAAAGAATTTC 780
Qy 781 AGCATGATTTCATATTGCTTCCACCAACTGCCAAATAACA 822
Db 781 AGCATGATTTCATATTGCTTCCACCAACTGCCAAATAACA 822

RESULT 4
US-09-898-883-1
; Sequence 1, Application US/09898883
; Patent No. 6585958
; GENERAL INFORMATION:
; APPLICANT: Little, Andrew Lamparski, Henry Schaur, Eric
; HENDERSON, Daniel
; TITLE OF INVENTION: ADENOVIRUS VECTORS SPECIFIC FOR CELLS EXPRESSING ALPHA-FETOPROTEIN AND METHODS OF USE THEREOF
; NUMBER OF SEQUENCES: 23
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: MORRISON & FORBSTER
; STREET: 755 PAGE MILL ROAD
; CITY: PALO ALTO
; STATE: CA
; ZIP: 94304-1018
; COUNTRY: USA
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/09/898, 883
; FILING DATE: 02-Jul-2001
; CLASSIFICATION: <Unknown>
; PRIORITY APPLICATION DATA:
; APPLICATION NUMBER: US/09/033, 428
; FILING DATE: <Unknown>
; ATTORNEY/AGENT INFORMATION:
; NAME: POLIZZI, CATHERINE M.
; REGISTRATION NUMBER: 40,130
; REFERENCE/DOCKET NUMBER: 34802-30004.00
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (415) 494-0792
; TELEFAX: (415) 494-0792
; INFORMATION FOR SEQ ID NO: 1:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 822 base pairs
; TYPE: nucleic acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; SEQUENCE DESCRIPTION: SEQ ID NO: 1:
US-09-898-883-1

Query Match 100.0%; Score 822; DB 4; Length 822;
 Best Local Similarity 100.0%; Pred. No. 1.5e-223; Length 822;
 Matches 822; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 GCATGGCTGGAACCTGTGACTTCTAGACATAAACATGAACTTGTGG 60
 Db 1 GCATGGCTGGAACCTGTGACTTCTAGACATAAACATGAACTTGTGG 60

Qy 61 GATTGTTGCTGTTAGCATACAAACTCTGGTCAAGCTCTCTTCAGCTTG 120
 Db 61 GATTGTTGCTGTTAGCATACAAACTCTGGTCAAGCTCTTCAGCTTG 120

Qy 121 AAAATTGGCTGTTCTCATGGTTCCTCTTCAGCTCATCTTCACACTCA 180
 Db 121 AAAATTGGCTGTTCTCATGGTTCCTCTTCAGCTCATCTTCACACTCA 180

Qy 181 CATGGCTACATAACTCTGGTCAAGCTTCAACTCTCATCTAGCTCANT 240
 Db 181 CATGGCTACATAACTCTGGTCAAGCTTCAACTCTCATCTAGCTCANT 240

Qy 241 CCTGGTCCAGAGATAAAAGTAGTATTCAAATGCACTTCAACTCTGGGCC 300
 Db 241 CCTGGTCCAGAGATAAAAGTAGTATTCAAATGCACTTCAACTCTGGGCC 300

Qy 301 TTAAAGACGTTCAACATACAACCGGGAGTTTGCCCTGGAATSTTCTTAATGCT 360
 Db 301 TTAAAGACGTTCAACATACAACCGGGAGTTTGCCCTGGAATSTTCTTAATGCT 360

Qy 361 CCTGTAAGCATAGGCTCCTCTGGTCTTAATCTTAATTACTTTAGCCACTGTCA 420
 Db 361 CCTGTAAGCATAGGCTCCTCTGGTCTTAATCTTAATTACTTTAGCCACTGTCA 420

Qy 421 TCCCACCTATGGGAGATGAGAGTAAAGGGAGCTGCTTAATTTAGCTAAGTCA 480
 Db 421 TCCCACCTATGGGAGATGAGAGTAAAGGGAGCTGCTTAATTTAGCTAAGTCA 480

Qy 481 TAGGCATAGGCCAGGACTGTTGGSTAACTGGTCACTTTATCTTAACTAATATTC 540
 Db 481 TAGGCATAGGCCAGGACTGTTGGSTAACTGGTCACTTTATCTTAACTAATATTC 540

Qy 541 CAAAATGAACTGACATGACTTGTACTAGCTTGTACTCTTCACTTCACTTCAG 600
 Db 541 CAAAATGAACTGACATGACTTGTACTAGCTTGTACTCTTCACTTCACTTCAG 600

Qy 601 CTTTATCCAGGCCACTTATAGCTCTGCTCTGAACTTAAACCGCTAT 660
 Db 601 CTTTATCCAGGCCACTTATAGCTCTGCTCTGAACTTAAACCGCTAT 660

Qy 661 GCTGTTAAATTATTGGCAAAATGCCATTTCACCTAACCTAACCTAACAGA 720
 Db 661 GCTGTTAAATTATTGGCAAAATGCCATTTCACCTAACCTAACCTAACAGA 720

Qy 721 TATACCAACAAAGGTTACTAGTTAACGGCATTCGCTGAAAGGATAAAAGATTTC 780
 Db 721 TATACCAACAAAGGTTACTAGTTAACGGCATTCGCTGAAAGGATAAAAGATTTC 780

Qy 781 AGCATGATTTCCATATTGGAAAGCTTCCACACTGCCAATAAACA 822
 Db 781 AGCATGATTTCCATATTGGAAAGCTTCCACACTGCCAATAAACA 822

RESULT 5 US-03-151-376-6
 ; Sequence 6, Application US/09151376
 ; Parent No. 6676935
 ; GENERAL INFORMATION:
 ; APPLICANT: Henderson, D.R.
 ; TITLE OF INVENTION: TISSUE SPECIFIC VIRAL VECTORS
 ; FILE REFERENCE: 348022000221
 ; CURRENT APPLICATION NUMBER: US/09/151,376
 ; CURRENT FILING DATE: 1998-09-10

; EARLIER APPLICATION NUMBER: 08/669,753
 ; EARLIER FILING DATE: 1995-06-26
 ; EARLIER APPLICATION NUMBER: 08/495,034
 ; EARLIER FILING DATE: 1995-06-27
 ; NUMBER OF SEQ ID NOS: 71
 ; SOFTWARE: PatentIn Ver. 2.0
 ; SEQ ID NO: 6
 ; LENGTH: 822
 ; TYPE: DNA
 ; ORGANISM: Homo sapiens
 ; US-09-151-376-6

Query Match 100.0%; Score 822; DB 4; Length 822;
 Best Local Similarity 100.0%; Pred. No. 1.5e-223; Length 822;
 Matches 822; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 GCATGGCTGTTCTGACTCTGGTCAAGCTCTCTTCAGCTCATCTTCACACTCA 180
 Db 1 GCATGGCTGTTCTGACTCTGGTCAAGCTCTCTTCAGCTCATCTTCACACTCA 180

Qy 61 GATTGTTGCTGTTAGCATACAAACTCTGGTCAAGCTCTTCAGCTCATCTTCACACTCA 180
 Db 61 GATTGTTGCTGTTAGCATACAAACTCTGGTCAAGCTCTTCAGCTCATCTTCACACTCA 180

Qy 121 AAAATTGGCTGTTCTCATGGTTCCTCTTCAGCTCATCTTCACACTCA 180
 Db 121 AAAATTGGCTGTTCTCATGGTTCCTCTTCAGCTCATCTTCACACTCA 180

Qy 181 CATGGCTACATAACTCTGGTCAAGCTTCAACTCTCATCTAGCTCANT 240
 Db 181 CATGGCTACATAACTCTGGTCAAGCTTCAACTCTCATCTAGCTCANT 240

Qy 241 CCTGGTCCAGAGATAAAAGTAGTATTCAAATGCACTTCAACTCTGGGCC 300
 Db 241 CCTGGTCCAGAGATAAAAGTAGTATTCAAATGCACTTCAACTCTGGGCC 300

Qy 301 TTAAAGACGTTCAACATACAACCGGGAGTTTGCCCTGGAATSTTCTTAATGCT 360
 Db 301 TTAAAGACGTTCAACATACAACCGGGAGTTTGCCCTGGAATSTTCTTAATGCT 360

Qy 361 CCTGTAAGCATAGGCCACTGTTGGSTAACTGGTCACTTTATCTTAACTAATATTC 420
 Db 361 CCTGTAAGCATAGGCCACTGTTGGSTAACTGGTCACTTTATCTTAACTAATATTC 420

Qy 421 CTTTCCAGAACATTAATGCTCTCTTCAGCTCATCTTCACACTCACTCTGGGGCC 300
 Db 421 CTTTCCAGAACATTAATGCTCTCTTCAGCTCATCTTCACACTCACTCTGGGGCC 300

Qy 481 TAGGCATAGGCCAGGACTGTTGGSTAACTGGTCACTTTATCTTAACTAATATTC 480
 Db 481 TAGGCATAGGCCAGGACTGTTGGSTAACTGGTCACTTTATCTTAACTAATATTC 480

Qy 541 CAAAATGAACTGACATGACTTGTACTAGCTTGTACTCTTCACTTCACTTCAG 540
 Db 541 CAAAATGAACTGACATGACTTGTACTAGCTTGTACTCTTCACTTCACTTCAG 540

Qy 601 CTTTATCCAGGCCACTTATAGCTCTGCTCTGAACTTAAACCGCTAT 660
 Db 601 CTTTATCCAGGCCACTTATAGCTCTGCTCTGAACTTAAACCGCTAT 660

Qy 661 GCTGTTAAATTATTGGCAAAATGCCATTTCACCTAACCTAACCTAACAGA 720
 Db 661 GCTGTTAAATTATTGGCAAAATGCCATTTCACCTAACCTAACCTAACAGA 720

Qy 721 TATACCAACAAAGGTTACTAGTTAACGGCATTCGCTGAAAGGATAAAAGATTTC 780
 Db 721 TATACCAACAAAGGTTACTAGTTAACGGCATTCGCTGAAAGGATAAAAGATTTC 780

Qy 781 AGCATGATTTCCATATTGGAAAGCTTCCACACTGCCAATAAACA 822
 Db 781 AGCATGATTTCCATATTGGAAAGCTTCCACACTGCCAATAAACA 822

Query Match 100.0%; Score 822; DB 4; Length 822;
 Best Local Similarity 100.0%; Pred. No. 1.5e-223; Length 822;
 Matches 822; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 GCATGGCTGTTCTGACTCTGGTCAAGCTCTCTTCAGCTCATCTTCACACTCA 180
 Db 1 GCATGGCTGTTCTGACTCTGGTCAAGCTCTCTTCAGCTCATCTTCACACTCA 180

Qy 61 GATTGTTGCTGTTAGCATACAAACTCTGGTCAAGCTCTTCAGCTCATCTTCACACTCA 180
 Db 61 GATTGTTGCTGTTAGCATACAAACTCTGGTCAAGCTCTTCAGCTCATCTTCACACTCA 180

Qy 121 AAAATTGGCTGTTCTCATGGTTCCTCTTCAGCTCATCTTCACACTCA 180
 Db 121 AAAATTGGCTGTTCTCATGGTTCCTCTTCAGCTCATCTTCACACTCA 180

Qy 181 CATGGCTACATAACTCTGGTCAAGCTTCAACTCTCATCTAGCTCANT 240
 Db 181 CATGGCTACATAACTCTGGTCAAGCTTCAACTCTCATCTAGCTCANT 240

Qy 241 CCTGGTCCAGAGATAAAAGTAGTATTCAAATGCACTTCAACTCTGGGCC 300
 Db 241 CCTGGTCCAGAGATAAAAGTAGTATTCAAATGCACTTCAACTCTGGGCC 300

Qy 301 TTAAAGACGTTCAACATACAACCGGGAGTTTGCCCTGGAATSTTCTTAATGCT 360
 Db 301 TTAAAGACGTTCAACATACAACCGGGAGTTTGCCCTGGAATSTTCTTAATGCT 360

Qy 361 CCTGTAAGCATAGGCCACTGTTGGSTAACTGGTCACTTTATCTTAACTAATATTC 420
 Db 361 CCTGTAAGCATAGGCCACTGTTGGSTAACTGGTCACTTTATCTTAACTAATATTC 420

Qy 421 CTTTCCAGAACATTAATGCTCTCTTCAGCTCATCTTCACACTCACTCTGGGGCC 300
 Db 421 CTTTCCAGAACATTAATGCTCTCTTCAGCTCATCTTCACACTCACTCTGGGGCC 300

Qy 481 TAGGCATAGGCCAGGACTGTTGGSTAACTGGTCACTTTATCTTAACTAATATTC 480
 Db 481 TAGGCATAGGCCAGGACTGTTGGSTAACTGGTCACTTTATCTTAACTAATATTC 480

Qy 541 CAAAATGAACTGACATGACTTGTACTAGCTTGTACTCTTCACTTCACTTCAG 540
 Db 541 CAAAATGAACTGACATGACTTGTACTAGCTTGTACTCTTCACTTCACTTCAG 540

Qy 601 CTTTATCCAGGCCACTTATAGCTCTGCTCTGAACTTAAACCGCTAT 660
 Db 601 CTTTATCCAGGCCACTTATAGCTCTGCTCTGAACTTAAACCGCTAT 660

Qy 661 GCTGTTAAATTATTGGCAAAATGCCATTTCACCTAACCTAACCTAACAGA 720
 Db 661 GCTGTTAAATTATTGGCAAAATGCCATTTCACCTAACCTAACCTAACAGA 720

Qy 721 TATACCAACAAAGGTTACTAGTTAACGGCATTCGCTGAAAGGATAAAAGATTTC 780
 Db 721 TATACCAACAAAGGTTACTAGTTAACGGCATTCGCTGAAAGGATAAAAGATTTC 780

Qy 781 AGCATGATTTCCATATTGGAAAGCTTCCACACTGCCAATAAACA 822
 Db 781 AGCATGATTTCCATATTGGAAAGCTTCCACACTGCCAATAAACA 822

RESULT 6
US-09-151-376-44
; Sequence 44, Application US/09151376
; Patent No. 6,676,935
; GENERAL INFORMATION
; APPLICANT: Henderson, D.R.
; APPLICANT: Schuur, E.R.
; TITLE OF INVENTION: TISSUE SPECIFIC VIRAL VECTORS
; FILE REFERENCE: 340022000221
; CURRENT FILING DATE: 1998-09-10
; EARLIER APPLICATION NUMBER: US/09/151,376
; EARLIER FILING DATE: 1996-06-26
; EARLIER APPLICATION NUMBER: 08/495,034
; EARLIER FILING DATE: 1995-06-27
; NUMBER OF SEQ ID NOS: 71
; SOFTWARE: PatentIn Ver. 2.0
; SBO ID NO. 44
; LENGTH: 822
; TYPE: DNA
; ORGANISM: Unknown
; FEATURE: Description of Unknown Organism: unknown
; OTHER INFORMATION: Description of Unknown Organism: unknown
US-09-151-376-44

Query Match 100.0%; Score 822; DB 4; Length 822;
Best Local Similarity 100.0%; Pred. No. 1.5e-223;
Matches 822; Conservative 0; Mismatches 0; Gaps 0;

Qy 1 GATTGGCTGGAATCTGTACTTGTAGGACTAAACTTTCAGCAATAACACATAGATTGAG 60
Db 1 GCATTGCTGTACTCTGTACTTGTAGGACTAAACTTTCAGCAATAACACATAGATTGAG 60

Qy 61 GATTGGTTGCTTTAGCTACAAACTCTGGTCAAACCTCCCTTTATGGCTTTGG 120
Db 61 GATTGGTTGCTTTAGCTACAAACTCTGGTCAAACCTCCCTTTATGGCTTTGG 120

Qy 121 AAAATTTCGTTCTCTCATGGTTCTCTTCACTGTATCTATTTCTAACCACTCA 180
Db 121 AAAATTTCGTTCTCTCATGGTTCTCTTCACTGTATCTATTTCTAACCACTCA 180

Qy 181 CATGGCTACAATAACTCTGGAAGGTATGATTCCTATCTAGCTCAT 240
Db 181 CATGGCTACAATAACTCTGCTGAAAGCTTCTATGATTCCTATCTAGCTCAT 240

Qy 241 CTGGTTCAAGAAGATAAAAGTAGTATTCAATGCACTCAACGTTCTCACCTGGGGGC 300
Db 241 CTGGTTCAAGAAGATAAAAGTAGTATTCAATGCACTCAACGTTCTCACCTGGGGC 300

Qy 301 TTAAAGAGCTTCAACATACAAACCGGGGAGTTTGCCTGAAATGTTCTAAAATGTT 360
Db 301 TTAAAGAGCTTCAACATACAAACCGGGGAGTTTGCCTGAAATGTTCTAAAATGTT 360

Qy 361 CCTGTAGCACATAGGTCTCTGTGTTCTAAATCTAAATCTACTTTCGCCAGTGCTCA 420
Db 361 CCTGTAGCACATAGGTCTCTGTGTTCTAAATCTAAATCTACTTTCGCCAGTGCTCA 420

Qy 421 TCCCACCTATGGGAGATGAGTGAAGAGGAGCTGTTCTAAATCAATAACCGCTAT 660
Db 421 TCCCACCTATGGGAGATGAGTGAAGAGGAGCTGTTCTAAATCAATAACCGCTAT 660

Qy 481 TAGGCATAGGCCAGGACTGTTGGTAACATGGTCACTTATCTAACTAAATAATTC 540
Db 481 TAGGCATAGGCCAGGACTGTTGGTAACATGGTCACTTATCTAACTAAATAATTC 540

Qy 541 CAAACTGAACTGTACTTAGTAACTGCTTGTACTTCACTTCAACACTCAG 600
Db 541 CAAACTGAACTGTACTTAGTAACTGCTTGTACTTCACTTCAACACTCAG 600

Qy 601 CTTTATCCAGGCCACTTATGACTCTGTCTGTTGAACTTAAATCAATAACCGCTAT 660
Db 601 CTTTATCCAGGCCACTTATGACTCTGTCTGTTGAACTTAAATCAATAACCGCTAT 660

Qy 661 GCTGTTAATTATGGCRAATGTCGCCATTTCACTTAAGGAATACTTAACTAAGTAACAGA 720
Db 661 GCTGTTAATTATGGCRAATGTCGCCATTTCACTTAAGGAATACTTAACTAAGTAACAGA 720

Qy 721 TATACCAACAAAGGTACTAGTTAACGGATGCTGAAAGGTATAAAAGAATTTC 780
Db 721 TATACCAACAAAGGTACTAGTTAACGGATGCTGAAAGGTATAAAAGAATTTC 780

Qy 781 ACCATGATTTCATATTGGCTTCCACCTGCTCCATAACA 822
Db 781 AGCATGATTTCATATTGGCTTCCACCTGCTCCATAACA 822

RESULT 7
US-09-814-351-8
; Sequence 8, Application US/09814351
; Patent No. 6632736
; GENERAL INFORMATION
; APPLICANT: Yu, De-Chao
; APPLICANT: Li, Yuanhao
; APPLICANT: Henderson, Daniel R.
; TITLE OF INVENTION: CELL-SPECIFIC ADENOVIRUS VECTORS
; TITLE OF INVENTION: COMPRISING AN INTERNAL RIBOSOME ENTRY SITE
; FILE REFERENCE: 348022601700
; CURRENT APPLICATION NUMBER: US/09/814,351
; PRIORITY FILING DATE: 2001-03-21
; PRIORITY APPLICATION NUMBER: US/09/814,351
; NUMBER OF SEQ ID NOS: 35
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO. 8
; LENGTH: 822
; TYPE: DNA
; ORGANISM: Artificial Sequence
; FEATURE: APB-TRE
; OTHER INFORMATION: APB-TRE
US-09-814-351-8

Query Match 100.0%; Score 822; DB 4; Length 822;
Best Local Similarity 100.0%; Pred. No. 1.5e-223;
Matches 822; Conservative 0; Mismatches 0; Gaps 0;

Qy 1 GCATTCTGTAACCTGTACTTGTAGGACTAAACTTGTAGCAATAACACATAGATTGAG 60
Db 1 GCATTCTGTAACCTGTACTTGTAGGACTAAACTTGTAGCAATAACACATAGATTGAG 60

Qy 121 AAAATTTCGTTCTCTCATGGTTCTCTTCACTGTATCTATTTCTAACACACTCA 180
Db 121 AAAATTTCGTTCTCTCATGGTTCTCTTCACTGTATCTATTTCTAACACACTCA 180

Qy 181 CATGGCTACAATAACTCTGGAAGGTATGATTCCTATCTAGCTCAT 240
Db 181 CATGGCTACAATAACTCTGCTGAAAGCTTCTATGATTCCTATCTAGCTCAT 240

Qy 241 CTGGTTCAAGAAGATAAAAGTAGTATTCAATGCACTCAACGTTCTCACCTGGGGGC 300
Db 241 CTGGTTCAAGAAGATAAAAGTAGTATTCAATGCACTCAACGTTCTCACCTGGGGC 300

Qy 301 TTAAAGAGCTTCAACATACAAACCGGGGAGTTTGCCTGAAATGTTCTAAAATGTT 360
Db 301 TTAAAGAGCTTCAACATACAAACCGGGGAGTTTGCCTGAAATGTTCTAAAATGTT 360

Qy 361 CCTGTAGCACATAGGTCTCTGTGTTCTAAATCTAAATCTACTTTCGCCAGTGCTCA 420
Db 361 CCTGTAGCACATAGGTCTCTGTGTTCTAAATCTAAATCTACTTTCGCCAGTGCTCA 420

Qy 421 TCCCACCTATGGGAGATGAGTGAAGAGGAGCTGTTCTAAATCAATAACCGCTAT 660
Db 421 TCCCACCTATGGGAGATGAGTGAAGAGGAGCTGTTCTAAATCAATAACCGCTAT 660

Qy 481 TAGGCATAGGCCAGGACTGTTGGTAACATGGTCACTTATCTAACTAAATAATTC 540
Db 481 TAGGCATAGGCCAGGACTGTTGGTAACATGGTCACTTATCTAACTAAATAATTC 540

Qy 541 CAAACTGAACTGTACTTAGTAACTGCTTGTACTTCACTTCAACACTCAG 600
Db 541 CAAACTGAACTGTACTTAGTAACTGCTTGTACTTCACTTCAACACTCAG 600

Qy 601 CTTTATCCAGGCCACTTATGACTCTGTCTGTTGAACTTAAATCAATAACCGCTAT 660
Db 601 CTTTATCCAGGCCACTTATGACTCTGTCTGTTGAACTTAAATCAATAACCGCTAT 660

Query Match 75.3%; Score 619; DB 1; Length 2240;
 Best Local Similarity 100.0%; Pred. No. 1e-165;
 Matches 619; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

1 GCAATGCTGTGAACCTGACTAACTGGATAACATAGATTGAG 1300
 481 TAGGCATAGGCCAGGACTGTTGGSTAAGTCACTTTATCTTAACATAATATTC 540
 481 TAGGCATAGGCCAGGACTGTTGGSTAAGTCACTTTATCTTAACATAATATTC 540
 541 CAAAAGTCACTGACATGACTTACTAGTTACTAAGSCTTGTGACTTACCATCCAG 600
 541 CAAAAGTCACTGACATGACTTACTAGTTACTAAGSCTTGTGACTTACCATCCAG 600
 601 CTTTATTCAGGCCACTTATAGCTCTGTGCTCTGAACTAAATAACCGTAT 660
 601 CTTTATTCAGGCCACTTATAGCTCTGTGCTCTGAACTAAATAACCGTAT 660
 601 GCTGTTAAATTATTGGCAAATGTCCTTCAACCTAACCTAAAGTAAACAGA 720
 601 GCTGTTAAATTATTGGCAAATGTCCTTCAACCTAACCTAAAGTAAACAGA 720
 721 TATACCCAAAGGTTTACAGGTTACAGGTTACCTGAAAGACTTAACAAATTC 780
 721 TATACCCAAAGGTTTACAGGTTACCTGAAAGACTTAACAAATTC 780
 781 AGCATGATTTCCAPATTGGCTTCCACCACTGGCCATATAACA 822
 781 AGCATGATTTCCAPATTGGCTTCCACCACTGGCCATATAACA 822

RESULT 8
 US-08-148-05BA-12
 Sequence 12, Application US/08148058A
 Patent No. 5804407

GENERAL INFORMATION:
 APPLICANT: TAMAOKI, TAIKI
 APPLICANT: NAKABAYASHI, HIDEKAZU
 TITLE OF INVENTION: IMPROVED METHOD OF EXPRESSING GENES IN
 TITLE OF INVENTION: MAMMALIAN CELLS
 NUMBER OF SEQUENCES: 46
 CORRESPONDENCE ADDRESS:
 ADDRESSEE: BURNS, DOANE, SWECKER & MATHIS
 STREET: 699 PRINCE STREET
 CITY: ALEXANDRIA
 STATE: VA
 COUNTRY: USA
 ZIP: 22313-1404

COMPUTER READABLE FORM:
 MEDIUM TYPE: Floppy disk
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: PatentIn Release #1.0, Version #1.25
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/08/148, 058A
 FILING DATE: 04-NOV-1993
 CLASSIFICATION: 435
 ATTORNEY/AGENT INFORMATION:
 NAME: MOOI, LESLIE A.
 REGISTRATION NUMBER: 37, 047
 REFERENCE/DOCKET NUMBER: 028722-074
 TELEPHONE: 415-854-8700
 TELEFAX: 415-854-8275
 INFORMATION FOR SEQ ID NO: 12:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 2240 base pairs
 TYPE: nucleic acid
 STRANDEDNESS: double
 TOPOLOGY: linear

US-08-148-05BA-12
 Query Match 75.3%; Score 619; DB 1; Length 2240;
 Best Local Similarity 100.0%; Pred. No. 1e-165;
 Matches 619; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

1 GCAATGCTGTGAACCTGACTAACTGGATAACATAGATTGAG 600
 61 GATTTTGGCTGTAGCATACAAACTCTGGTCAAAAGCTCTCTTATTGCTTGG 120
 61 GATTTTGGCTGTAGCATACAAACTCTGGTCAAAAGCTCTCTTATTGCTTGG 120
 1302 GATTTTGGCTGTAGCATACAAACTCTGGTCAAAAGCTCTCTTATTGCTTGG 136
 121 AAAATTGGCTGTTCTCATGGTTCTCATGCTATCTATTATTTCTAACACTCA 180
 121 AAAATTGGCTGTTCTCATGGTTCTCATGCTATCTATTATTTCTAACACTCA 180
 1362 AAAATTGGCTGTTCTCATGGTTCTCATGCTATCTATTATTTCTAACACTCA 142
 181 CATGGCTACATAACTGTCCTGCAACGTTATGATCCTAGCTCAAT 240
 181 CATGGCTACATAACTGTCCTGCAACGTTATGATCCTAGCTCAAT 240
 1422 CATGGCTACATAACTGTCCTGCAACGTTATGATCCTAGCTCAAT 148
 241 CTTGTTCCAGAAAGATAAAAGTAGTATTCAATGGACATCAACGCTCCACTGGGGC 300
 241 CTTGTTCCAGAAAGATAAAAGTAGTATTCAATGGACATCAACGCTCCACTGGGGC 300
 1482 CTTGTTCCAGAAAGATAAAAGTAGTATTCAATGGACATCAACGCTCCACTGGGGC 154
 301 TTAAAGACGTTACACATAAAAGGGAGTTGGCTTGAATTTCTAAATGTGT 360
 301 TTAAAGACGTTACACATAAAAGGGAGTTGGCTTGAATTTCTAAATGTGT 360
 1542 TTAAGACGTTACACATAAAAGGGAGTTGGCTTGAATTTCTAAATGTGT 160
 1601 CCTGTAGCACATGGCTCCCTTGTCTTAAACTTAATCTTAAGCTGCTCA 420
 1601 CCTGTAGCACATGGCTCCCTTGTCTTAAACTTAATCTTAAGCTGCTCA 420
 1602 CCTGTAGCACATGGCTCCCTTGTCTTAAACTTAATCTTAAGCTGCTCA 166
 421 TCCCCCCTATGGGAGATGAGCTAACGGAGCTGTTCTTAAACTTAACCTAGTCAA 480
 421 TCCCCCCTATGGGAGATGAGCTAACGGAGCTGTTCTTAAACTTAACCTAGTCAA 480
 1662 TCCCCCCTATGGGAGATGAGCTAACGGAGCTGTTCTTAAACTTAACCTAGTCAA 172
 481 TAGGGATAGGCCAGGACTGTTGGTAAACTGGGACTTAACTAAATATATC 540
 481 TAGGGATAGGCCAGGACTGTTGGTAAACTGGGACTTAACTAAATATATC 540
 1722 TAGGGATAGGCCAGGACTGTTGGTAAACTGGGACTTAACTAAATATATC 178
 541 CAAACTGAACTGACTGACTTAACTGACTTAACTGACTTAACTGACTTAACTGACTCAG 600
 541 CAAACTGAACTGACTGACTTAACTGACTTAACTGACTTAACTGACTTAACTGACTCAG 600
 1782 CAAACTGAACTGACTGACTTAACTGACTTAACTGACTTAACTGACTTAACTGACTCAG 184
 601 CTTTATCCAGGCCACTTAT 619
 601 CTTTATCCAGGCCACTTAT 619
 1842 CTTTATCCAGGCCACTTAT 1860
 1842 CTTTATCCAGGCCACTTAT 1860

RESULT 9
 US-08-478-042-12
 Sequence 12, Application US/08478042
 ; Patent No. 5807738

GENERAL INFORMATION:
 APPLICANT: TAMAOKI, TAIKI
 APPLICANT: NAKABAYASHI, HIDEKAZU
 TITLE OF INVENTION: IMPROVED METHOD OF EXPRESSING GENES IN
 TITLE OF INVENTION: MAMMALIAN CELLS
 NUMBER OF SEQUENCES: 46
 CORRESPONDENCE ADDRESS:
 ADDRESSEE: BURNS, DOANE, SWECKER & MATHIS
 STREET: 699 PRINCE STREET
 CITY: ALEXANDRIA
 STATE: VA
 COUNTRY: USA
 ZIP: 22313-1404
 COMPUTER READABLE FORM:
 MEDIUM TYPE: Floppy disk
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: PatentIn Release #1.0, Version #1.25
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/08/148, 058A
 FILING DATE: 04-NOV-1993
 CLASSIFICATION: 435
 ATTORNEY/AGENT INFORMATION:
 NAME: MOOI, LESLIE A.
 REGISTRATION NUMBER: 37, 047
 REFERENCE/DOCKET NUMBER: 028722-074
 TELEPHONE: 415-854-8700
 TELEFAX: 415-854-8275
 INFORMATION FOR SEQ ID NO: 12:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 2240 base pairs
 TYPE: nucleic acid
 STRANDEDNESS: double
 TOPOLOGY: linear

US-08-478-042-12
 COMPUTER READABLE FORM:
 MEDIUM TYPE: Floppy disk
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: PatentIn Release #1.0, Version #1.25
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/08/478, 042
 FILING DATE: 07-JUN-1995
 CLASSIFICATION: 435
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: US 08/148, 058
 FILING DATE: 04-NOV-1993
 ATTORNEY/AGENT INFORMATION:

NAME : MOOI, LESLIE A. ;
 REGISTRATION NUMBER : 37,047 ;
 REFERENCE/DOCKET NUMBER : 028722-126 ;
 TELECOMMUNICATION INFORMATION :
 TELEPHONE : 415-854-7400 ;
 TELEFAX : 415-854-8275 ;
 INFORMATION FOR SEQ ID NO: 12:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 2240 base pairs
 TYPE: nucleic acid
 STRANDEDNESS: double
 TOPOLOGY: linear
 US - 08-478-042-12

| | Query Match | Score | Length | DB | Indels | Gaps | No. 1e-165; | Mismatches | Best Local Similarity | Length | DB 1; | Length 2240; |
|----|-------------|---|--------|--------|--------|------|-------------|------------|-----------------------|--------|-------|--------------|
| | Matches | 619; | 619; | 100 %; | Pred. | 0; | ; | 0; | ; | 2240 | ; | ; |
| Qy | 1 | GCATTCGTGAACTGTACTTGGACTAAACCTGGATAAACACATAGATTGAG | 60 | | | | | | | | | |
| Db | 1242 | GCATTCGTGAACTGTACTTGGACTAAACCTGGATAAACACATAGATTGAG | 1301 | | | | | | | | | |
| Qy | 61 | GATTGTTGCTGTTGACATAGAACACTCGGTTCAAGCTCCTCTTATTGCTGCTTGG | 120 | | | | | | | | | |
| Db | 1302 | GATTGTTGCTGTTGACATAGAACACTCGGTTCAAGCTCCTCTTATTGCTGCTTGG | 1361 | | | | | | | | | |
| Qy | 121 | AAAATTGCGTGTCTCATGGTTCTTCACTGCTCTATCTTCTAACCCATCTCA | 180 | | | | | | | | | |
| Db | 1362 | AAAATTGCGTGTCTCATGGTTCTTCACTGCTCTATCTTCTAACCCATCTCA | 1421 | | | | | | | | | |
| Qy | 181 | CATGGCTACATAACTGTCAGAGTTGATTCTCAAAATATCTAGCCCTCAAT | 240 | | | | | | | | | |
| Db | 1422 | CATGGCTACATAACTGTCAGAGTTGATTCTCAAAATATCTAGCCCTCAAT | 1481 | | | | | | | | | |
| Qy | 241 | CTTGTTCAGAAGATAAAAGTAGTTCAATGCAACCTCTCCACCTGGAGGGC | 300 | | | | | | | | | |
| Db | 1482 | CTTGTTCAGAAGATAAAAGTAGTTCAATGCAACCTCTCCACCTGGAGGGC | 1541 | | | | | | | | | |
| Qy | 301 | TTRAGACGTITCACTACATAGAACCGGGAGTTGCCCTTAAATCTTAATGTT | 360 | | | | | | | | | |
| Db | 1542 | TTRAGACGTITCACTACATAGAACCGGGAGTTGCCCTTAAATCTTAATGTT | 1601 | | | | | | | | | |
| Qy | 361 | CCTGTTAGCACTAGGGGAGATGAGAGTGAAGGGAGCTGATTAATTTACACTA | 480 | | | | | | | | | |
| Db | 1602 | CCTGTTAGCACTAGGGGAGATGAGAGTGAAGGGAGCTGATTAATTTACACTA | 1661 | | | | | | | | | |
| Qy | 421 | TCCCACCTATGGGAGATGAGAGTGAAGGGAGCTGATTAATTTACACTA | 480 | | | | | | | | | |
| Db | 1662 | TCCCACCTATGGGAGATGAGAGTGAAGGGAGCTGATTAATTTACACTA | 1721 | | | | | | | | | |
| Qy | 481 | TAGGGTAGGGCAGGACTGTTGGGTAACCTGTACTTCTTAATCTAACTATTC | 540 | | | | | | | | | |
| Db | 1722 | TAGGGTAGGGCAGGACTGTTGGGTAACCTGTACTTCTTAATCTAACTATTC | 1781 | | | | | | | | | |
| Qy | 541 | CAAAACTGACATGTTACTAAGTCTTGTACTTCTTGTACTTCTTGTACTTCT | 600 | | | | | | | | | |
| Db | 1782 | CAAACTGACATGTTACTAAGTCTTGTACTTCTTGTACTTCTTGTACTTCT | 1841 | | | | | | | | | |
| Qy | 601 | CTTATCCAGGCCACTTAT | 619 | | | | | | | | | |
| Db | 1842 | CTTATCCAGGCCACTTAT | 1860 | | | | | | | | | |

RESULT 10

US - 08-645-215-12

Sequence 12, Application US/08645215

; Patent No. 58276616

GENERAL INFORMATION:

; APPLICANT: NAKABAYASHI, TAIKI

; ADDRESS: HIDEKAZU

TITLE OF INVENTION: IMPROVED METHOD OF EXPRESSING GENES IN

; TITLE OF INVENTION: MAMMALIAN CELLS

; NUMBER OF SEQUENCES: 46

Qy 541 CAAACTGACATGTACTTGTACTAAGTTGTTGACTTATCATTCAACCACTCG 600
 Db 1782 CAAACTGACATGTACTTGTACTAAGTTGTTGACTTATCATTCAACCACTCG 1841

Qy 601 CTTATCCAGGCCACTTAT 619
 Db 1842 CTTATCCAGGCCACTTAT 1860

RESULT 1.1
 US-08-466-604-12
 Sequence 12, Application US/08466604
 Patent No. 5843776
 GENERAL INFORMATION:
 APPLICANT: TAMAOKI, TAIKI
 ADDRESS: NAKABAYASHI, HIDENAKU
 TITLE OF INVENTION: IMPROVED METHOD OF EXPRESSING GENES IN MAMMALIAN CELLS
 NUMBER OF SEQUENCES: 46
 STATE/CITY: ALEXANDRIA
 COUNTRY: USA
 ZIP: 22313-1404
 COMPUTER READABLE FORM:
 MEDIUM TYPE: Floppy disk
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: PatentIn Release #1.0, Version #1.25
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/08/466,604
 FILING DATE: 06-JUN-1995
 CLASSIFICATION: 435
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: US 08/148,058
 FILING DATE: 04-NOV-1993
 ATTORNEY/AGENT INFORMATION:
 NAME: MOOL, LESLIE A.
 REGISTRATION NUMBER: 37,047
 REFERENCE/DOCKET NUMBER: Q28722-125
 TELECOMMUNICATION INFORMATION:
 TELEPHONE: 415-554-7400
 TELEFAX: 415-854-8275
 INFORMATION FOR SEQ ID NO: 12:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 2240 base pairs
 TYPE: nucleic acid
 STRANDEDNESS: double
 TOPOLOGY: linear

US-08-466-604-12

Query Match 75.3%; Score 619; DB 2; length 2240;
 Best Local Similarity 100.0%; Pred. No 1e-165;
 Matches 619; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 GCATGGCTGTGAACCTGTGTTACTTAGGACTAAACTTGAGGATAACACATAGATGAG 60
 Db 1242 GCATGGCTGTGAACCTGTGTTACTTAGGACTAAACTTGAGGATAACACATAGATGAG 1301

Qy 61 GATTTGGTTGGTTGAGCATCAAACCTGGTTCAALAGTCCTCTTGTGG 120
 Db 1302 GATTTGGTTGGTTGAGCATCAAACCTGGTTCAALAGTCCTCTTGTGG 1361

Qy 121 AAAATTGGTGTCTTCATGGTTCTCTCTTCACTGCTATCTATTTCTCAACCACTCA 180
 Db 1362 AAAATTGGTGTCTTCATGGTTCTCTCTTCACTGCTATCTATTTCTCAACCACTCA 1421

Qy 181 CATGGCTACATAACTGTGTCGAACGCTATGATTCGCCAATATCTATCTAGCTCTCAAT 240
 Db 1422 CATGGCTACATAACTGTGTCGAACGCTATGATTCGCCAATATCTATCTAGCTCTCAAT 1481

Qy 241 CTGTGTCAGAAGATAAAAGTAGTATCAATGCACTAACGTCCTCCACTGGGGGC 300
 Db 1482 CTGTGTCAGAAGATAAAAGTAGTATCAATGCACTAACGTCCTCCACTGGGGC 1541

Qy 301 TAAAGACGTTCAACATACAAACGGGAGTTGCTCTGAATTTCTTAAATGTGT 360
 Db 1542 TAAAGACGTTCAACATACAAACGGGAGTTGCTCTGAATTTCTTAAATGTGT 1600

Qy 361 CCTGTAACATAGGTGCTCTCTTAAATCTAAATTACTTTAGGCCAGTGCTCA 420
 Db 1602 CCTGTAACATAGGTGCTCTCTTAAATCTAAATTACTTTAGGCCAGTGCTCA 1666

Qy 421 TCCACCTATGGGAGATGAGGTGAAAGGGGCTGATTAATTAATTAGTCAA 480
 Db 1662 TCCACCTATGGGAGATGAGGTGAAAGGGGCTGATTAATTAATTAGTCAA 1722

Qy 481 TAGCATAGGCCAGACTGTTGGTAAACTCTTAAACTAAATATTC 540
 Db 1722 TAGCATAGGCCAGACTGTTGGTAAACTCTTAAACTAAATATTC 1784

Qy 541 CAAACATGAACTGTACTTGTACTAAGTCTTGACTTTATCATTCAACCTCG 600
 Db 1782 CAAACATGAACTGTACTTGTACTAAGTCTTGACTTTATCATTCAACCTCG 184

Qy 601 CTTTATCCAGGCCACTTAT 619
 Db 1842 CTTTATCCAGGCCACTTAT 1860

RESULT 1.2
 US-09-033-428-2
 Sequence 2, Application US/09033428
 Patent No. 6254962
 GENERAL INFORMATION:
 APPLICANT: Little, Andrew
 APPLICANT: Lamparski, Henry
 APPLICANT: Schaur, Eric
 APPLICANT: Henderson, Daniel
 TITLE OF INVENTION: ADENOVIRUS VECTORS SPECIFIC FOR CELLS
 TITLE OF INVENTION: EXPRESSING ALPHA-FETOPROTEIN AND METHODS OF USE THEREOF
 NUMBER OF SEQUENCES: 23
 CORRESPONDENCE ADDRESS:
 ADDRESSEE: MORRISON & FOERSTER
 STREET: 755 PAGE MILL ROAD
 CITY: PALO ALTO
 STATE: CA
 COUNTRY: USA
 ZIP: 94104-1018
 COMPUTER READABLE FORM:
 MEDIUM TYPE: Floppy disk
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: Patent In Release #1.0, Version #1.30
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/09/033,428
 FILING DATE: 04-NOV-1993
 ATTORNEY/AGENT INFORMATION:
 NAME: POLIZZI, CATHERINE M.
 REFERENCE/DOCKET NUMBER: 34802-30004.00
 TELECOMMUNICATION INFORMATION:
 TELEPHONE: (415) 813-5600
 TELEFAX: (415) 494-0792
 TELEX: 706141 MRSINFOERS SFO
 INFORMATION FOR SEQ ID NO: 2:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 5224 base pairs
 TYPE: nucleic acid
 STRANDEDNESS: double
 TOPOLOGY: linear

US-09-033-428-2

Query Match 75.3%; Score 61.9; DB 3; Length 5224;
 Best Local Similarity 100.0%; Pred. No. 1.4e-165;
 Matches 619; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 GCATTGCTGTGAACCTCTGACTTCTAGGACTAAACAGCAATAGATTGAG 60
 Db 1242 GCATTGCTGTGAACCTCTGACTTCTAGGACTAAACAGCAATAGATTGAG 1301

Qy 61 GATTGTRGCTGTAGCATACAAACTCTGGTCAAAGGCCCTTTATGCCTGTCCTGG 120
 Db 1302 GATTGTRGCTGTAGCATACAAACTCTGGTCAAAGGCCCTTTATGCCTGTCCTGG 1361

Qy 121 AAATTCTCTGTCTCTGTTCTCTTTTCACTGCTATCPATTCTCACCTCA 180
 Db 1362 AAATTCTGTGTTCTCTGTTCTCTTTTCACTGCTATCPATTCTCACCTCA 1421

Qy 181 CATTGCTACAATAAATCTGTGAGCTPATGATTCCAAATATCTATCTAGCCTCAT 240
 Db 1422 CATTGCTACAATAAATCTGTGAGCTPATGATTCCAAATATCTATCTAGCCTCAT 1481

Qy 241 CTGTTCCAGAGATAAAAAGTAGTATTCAAAATGCACTCAAGTCAGTCCTCACTGGGGC 300
 Db 1482 CTGTTCCAGAGATAAAAAGTAGTATTCAAAATGCACTCAAGTCAGTCCTCACTGGGGC 1541

Qy 301 TTAAGAGCTTCAACATACAAACCGGGAGTTGGCTGGATGTTCTAAATGTT 360
 Db 1542 TTAAGAGCTTCAACATACAAACCGGGAGTTGGCTGGATGTTCTAAATGTT 1601

Qy 361 CCTGAGCACATAGGTGCTCTGTCCTAAATCTAATTACTTTAGCCAGTGCTCA 420
 Db 1602 CCTGAGCACATAGGTGCTCTGTCCTAAATCTAATTACTTTAGCCAGTGCTCA 1661

Qy 421 TCCCACCATGGGAGATGGAGTGAAGGGAGCCCTGTTAAATACAACTCAA 480
 Db 1662 TCCCACCATGGGAGATGGAGTGAAGGGAGCCCTGTTAAATACAACTCAA 1721

Qy 481 TAGGCATAGGCCAGGACTGTTGGTAACACTGTCCTTATCTAACTAAATAATTC 540
 Db 1722 TAGGCATAGGCCAGGACTGTTGGTAACACTGTCCTTATCTAACTAAATAATTC 1781

Qy 541 CAAACTGACATGTGACTTAGTACTAGTACTAGTCTTGTACTTAACTACCTCAG 600
 Db 1782 CAAACTGACATGTGACTTAGTACTAGTACTAGTCTTGTACTTAACTACCTCAG 1841

Qy 601 CTTTATCAGGCCACTTAT 61.9
 Db 1642 CTTTATCAGGCCACTTAT 1860

RESULT 13
 US-09-033-556-5
 Sequence 5, Application US/09033556
 Patent No. 6412700
 GENERAL INFORMATION:
 APPLICANT: Henderson, Daniel R.
 APPLICANT: Yu, De chao
 TITLE OF INVENTION: ADENOVIRUS VECTORS CONTAINING
 HETEROLOGOUS TRANSCRIPTION REGULATORY ELEMENTS AND METHODS
 TITLE OF INVENTION: OF USING SAME
 NUMBER OF SEQUENCES: 41
 CORRESPONDENCE ADDRESS:
 ADDRESSEE: MORRISON & FOERSTER
 STREET: 755 PAGE MILL ROAD
 CITY: Palo Alto
 STATE: CA
 COUNTRY: USA
 ZIP: 94104-1018
 COMPUTER READABLE FORM:
 MEDIUM TYPE: Diskette
 COMPUTER: IBM Compatible
 OPERATING SYSTEM: Windows
 SOFTWARE: FASTSEQ for Windows Version 2.0b

Qy 1601 CTTTATCAGGCCACTTAT 61.9
 Db 1842 CTTTATCAGGCCACTTAT 1860

RESULT 14

US-09-474-699-15
Sequence 15, Application US/09474699
PATENT NO. 6585968
GENERAL INFORMATION:
APPLICANT: Henderson, Daniel R.
APPLICANT: Yu, De Chao
TITLE OF INVENTION: TARGET CELL-SPECIFIC ADENOVIRAL VECTORS
FILE REFERENCE: 34802201300
CURRENT APPLICATION NUMBER: US/09/474,699
CURRENT FILING DATE: 1999-12-29
PRIOR APPLICATION NUMBER: 60/114,262
PRIOR FILING DATE: 1998-12-30
NUMBER OF SEQ ID NOS: 23
SEQ ID NO: 15
LENGTH: 5224
TYPE: DNA
ORGANISM: Homo Sapien
US-09-474-699-15

Query Match 75.3%; Score 619; DB 4; Length 5224;
Best Local Similarity 100.0%; Pred. No. 1 4e-165;
Matches 619; Conservative 0; Mismatches 0; Gaps 0;

| | | |
|----|------|--|
| Qy | 1 | GCAATGCTGAAACTGTGACTTGGAACTAACACATAGATTGAG 60 |
| Db | 1242 | GCATGCTGAACTGTGACTTGGAACTAACATAGATTGAG 1301 |
| Qy | 61 | GATTGTTGCTGTAGCATACAACACTGGTTCAAGCTCTCTTATGCTTGCTTG 120 |
| Db | 1302 | GATTGTTGCTGTAGCATACAACACTGGTTCAAGCTCTCTTATGCTTGCTTG 1361 |
| Qy | 121 | AAAATTGCTGTTCTCATGGTTCTCATGGTTCTCATGGCTCAACACATGATTGAG 180 |
| Db | 1362 | AAAATTGCTGTTCTCATGGTTCTCATGGTTCTCATGGCTCAACACATGATTGAG 1421 |
| Qy | 181 | CATGGCTACAATAACTGTGCAAGGTTATGATTCCCAAAATATCATCTAGCTCAAT 240 |
| Db | 1422 | CATGGCTACAATAACTGTGCAAGGTTATGATTCCCAAAATATCATCTAGCTCAAT 1481 |
| Qy | 241 | CTTGTTCAGAGATAAAAGTAGATTCAAATGCAACGCTCACAATGGGCC 300 |
| Db | 1482 | CTTGTTCAGAGATAAAAGTAGATTCAAATGCAACGCTCACAATGGGCC 1541 |
| Qy | 301 | TAAAGAGTTCAAGATAACAACCGGGAGTTGCCGAATTTCTAAATGTT 360 |
| Db | 1542 | TAAAGAGTTCAAGATAACAACGGGAGTTGCCGAATTTCTAAATGTT 1601 |
| Qy | 361 | CCTGTAGCACATAGGGTCCTCTGGCTTAAATCTTAATTACTTGGCCAGTGCTCA 420 |
| Db | 1602 | CCTGTAGCACATAGGGTCCTCTGGCTTAAATCTTAATTACTTGGCCAGTGCTCA 1661 |
| Qy | 421 | TCCCACCTATGGGAGATGAGAGTAAAGGGAGCTGTTAATTAACATAAGTCAA 480 |
| Db | 1662 | TCCCACCTATGGGAGATGAGAGTAAAGGGAGCTGTTAATTAACATAAGTCAA 1781 |
| Qy | 541 | CAAAACTGAACTGATGACTTAACTAAGTCTTGGACTTATCTCATACCACTAG 600 |
| Db | 1782 | CAAAACTGAACTGATGACTTAACTAAGTCTTGGACTTATCTCATACCACTAG 1841 |
| Qy | 601 | CTTATCCAGGCCACTTAT 619 |
| Db | 1842 | CTTATCCAGGCCACTTAT 1860 |

Query Match 75.3%; Score 619; DB 4; Length 5224;
Best Local Similarity 100.0%; Pred. No. 1 4e-165;
Matches 619; Conservative 0; Mismatches 0; Gaps 0;

| | | |
|----|------|--|
| Qy | 1 | GCATTGCTGTAACCTGTGACTCTGTGACTTGGAAATACACATAGATTGAG 60 |
| Db | 1242 | GCATTGCTGTAACCTGTGACTCTGTGACTTGGAAATACACATAGATTGAG 1301 |
| Qy | 61 | GATTGTTGCTGTAGCATACAACACTGGTTCAAGCTCTCTTATGCTTGCTTG 120 |
| Db | 1302 | GATTGTTGCTGTAGCATACAACACTGGTTCAAGCTCTCTTATGCTTGCTTG 1361 |
| Qy | 121 | AAAATTGCTGTTCTCATGGTTCTCATGGTTCTCATGGCTCAACACATGATTGAG 180 |
| Db | 1362 | AAAATTGCTGTTCTCATGGTTCTCATGGTTCTCATGGCTCAACACATGATTGAG 1421 |
| Qy | 181 | CATGGCTACAATAACTGTGCAAGGTTATGATTCCCAAAATATCATCTAGCTCAAT 240 |
| Db | 1422 | CATGGCTACAATAACTGTGCAAGGTTATGATTCCCAAAATATCATCTAGCTCAAT 1481 |
| Qy | 241 | CTTGTTCAGAGATAAAAGTAGATTCAAATGCAACGCTCACAATGGGCC 300 |
| Db | 1542 | CTTGTTCAGAGATAAAAGTAGATTCAAATGCAACGCTCACAATGGGCC 1541 |

RESULT 15
US-09-898-883-2 Application US/09888883

RESULT 2 Application US/09888883

| | | | |
|----|------|--|------|
| Qy | 361 | CCTGTAGCACATAGGGTCCCTCTGTTCTAAATCTAATTACTTTAGCCCCAGTGCTCA | 420 |
| Db | 1602 | CCTGTAGCACATAGGGTCCCTCTGTTCTAAATCTAATTACTTTAGCCCCAGTGCTCA | 1661 |
| Qy | 421 | TCCACCATGCGGAGTGTAGAGTGTAAAGGGAGCCGTTAAATTAATCTAATTACTTTAGCCCCAGTGCTCA | 480 |
| Db | 1662 | TCCACCATGCGGAGTGTAGAGTGTAAAGGGAGCCGTTAAATTAATCTAATTACTTGCTCA | 1721 |
| Qy | 481 | TAGGCATAGGCCAGGACTGTTGGTAAACTCTGTTCACTTTATCTTAACCTAAATAATATTC | 540 |
| Db | 1722 | TAGGCATAGGCCAGGACTGTTGGTAAACTCTGTTCACTTTATCTTAACCTAAATAATATTC | 1781 |
| Qy | 541 | CAAACCTGAACATGTACTTGTACTTAAGTCTTGACTTTATCTCATTCACTACCTCAG | 600 |
| Db | 1782 | CAAACCTGAACATGTACTTGTACTTAAGTCTTGACTTTATCTCATTCACTACCTCAG | 1841 |
| Qy | 601 | CTTTATCCAGGCCACTTAT | 619 |
| Db | 1842 | CTTTATCCAGGCCACTTAT | 1860 |

Search completed: August 6, 2005, 23:02:37
Job time : 130.256 sec

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